### 346/2825 FIGURE 318

TCTACGAGAGCCTCCTGTCGCTGAGCCCTGACGTGCCCGTGCCATCCGACCATGGAGGGACTGAGTCCAGCCCAG  $\tt CCCGCCTGGGCCCTGAGCTGTCACCCTCACCCACTTCGCCCACTGCAACCTCCACCACCCCCTCGCGCTACAAGA$ CTGAGCTATGTCGGACCTTCTCAGAGAGTGGGCGCTGCCGCTACGGGGCCAAGTGCCAGTTTGCCCATGGCCTGG GCGAGCTGCGCCAGGCCAATCGCCACCCCAAATACAAGACGGAACTCTGTCACAAGTTCTACCTCCAGGGCCGCT GCCCTACGGCTCTCGCTGCCACTTCATCCACAACCCTAGCGAAGACCTGGCGGCCCCGGGCCACCCTCCTGTGC CTTCCCTGTCCTCCAGCTCCTTCTCGCCCTCCAGCTCCCACCACCACCTGGGGACCTTCCACTGTCACCCTCTG CCTTCTCTGCTGCCCCTGGCACCCCCTGGCTCGAAGAGACCCCCACCCCAGTCTGTTGCCCCTCCTGCCGAAGGG CTGATGAATATGCCAGCAGCGGCAGCCTGGGGGGGCTCTGACTCTCCCGTCTTCGAGGCGGGAGTTTTTGCAC CACCCAGCCCGTGGCAGCCCCCCGGCGACTCCCCATCTTCAATCGCATCTCTGTTTCTGAG**TGA**CAAAGTGACT GCCCGGTCAGATCAGCTGGATCTCAGCGGGGAGCCACGTCTCTTGCACTGTGGTCTCTGCATGGACCCCAGGGCT GTGGGGACTTGGGGGACAGTAATCAAGTAATCCCCTTTTCCAGAATGCATTAACCCACTCCCCTGACCTCACGCT GGGGCAGGTCCCCAAGTGTGCAAGCTCAGTATTCATGATGGTGGGGGATGGAGTGTCTTCCGAGGTTCTTGGGGG AGAATCTTATGTGCTGTGAATAATAGGCCTTCACTGCCCCTCCAGTTTTTATAGACCTGAGGTTCCAGTGTCTCC TGGTAACTGGAACCTCTCCTGAGGGGGAATCCTGGTGCTCAAATTACCCTCCAAAAGCAAGTAGCCAAAGCCGTT TTATATATTGGGTCGTCTCCCTTGTATTTTCTTCCTTTTTTTGTAATATTGAAAACGACGATATAATT ATTATAAGTAGACTATAATATTATTAGTAATATATTATTATTACCTTAAAAGTCTATTTTTGTGTTTTTGGGCATTT TTAAATAAACAATCTGAGTGT

# 347/2825 FIGURE 319

MDLTAIYESLLSLSPDVPVPSDHGGTESSPGWGSSGPWSLSPSDSSPSGVTSRLPGRSTSLVEGRSCGWVPPPPG FAPLAPRLGPELSPSPTSPTATSTTPSRYKTELCRTFSESGRCRYGAKCQFAHGLGELRQANRHPKYKTELCHKF YLQGRCPYGSRCHFIHNPSEDLAAPGHPPVLRQSISFSGLPSGRRTSPPPPGLAGPSLSSSSFSPSSSPPPPGDL PLSPSAFSAAPGTPLARRDPTPVCCPSCRRATPISVWGPLGGLVRTPSVQSLGSDPDEYASSGSSLGGSDSPVFE AGVFAPPQPVAAPRRLPIFNRISVSE

### 348/2825 FIGURE 320

# 349/2825 FIGURE 321

MSSIGTGYDLSASTFSPDGRVFQVEYAMKAVENSSTAIGIRCKDGVVFGVEKLVLSKLYEEGSNKRLFNVDRHVG MAVAGLLADARSLADIAREEASNFRSNFGYNIPLKHLADRVAMYVHAYTLYSAVRPFGCSFMLGSYSVNDGAQLY MIDPSGVSYGYWGCAIGKARQAAKTEIEKLQMKEMTCRDIVKEVAKIIYIVHDEVKDKAFELELSWVGELTNGRH EIVPKDIREEAEKYAKESLKEEDESDDDNM

### 350/2825 FIGURE 322

GCACGAGGCTCCGCCTCGCTTGCTCCTGCCGGGCGTGCAGGCCCCGCCGCCGCCATGTCGGGCTCGTTCGAGCT GGTCACCCCGCGGATCTACGTGGGCAACGCGTCTGTGGCTCAGGACATCCCCAAGCTGCAGAAACTAGGCATCAC CCATGTGCTGAACGCGGCTGAGGGCAGGTCCTTCATGCACGTCAACACCAATGCCAACTTCTACAAGGACTCCGG CATCACATACCTGGGCATCAAGGCCAACGACACACAGGAGTTCAACCTCAGCGCTTACTTTGAAAGGGCTGCCGA CTTCATTGACCAGGCTTTGGCTCAAAAGAATGGCCGGGTGCTCGTCCACTGCCGGGAAGGTTATAGCCGCTCCCC AACGCTAGTTATCGCCTACCTCATGATGCGGCAGAAGATGGACGTCAAGTCTGCCCTGAGCATCGTGAGGCAGAA CCGTGAGATCGGCCCAACGATGGCTTCCTGGCCAGCTCTGCCAGCTCAATGACAGACTAGCCAAGGAGGGGGAA GCTGCCATGTTTAGGAAACACACTGTACCCTGCTCCCAGCATCACAAGGCACTTGTCTACAAGTGTGTCCCAACA CAGTCCTGGGCCACTTTCCCCACCCTGGGGAGCACATAAAGAAGCTTGCCAAGGGGGGGCGTCCTTGCTCCCCAGT TGTCCTGTTTCTGTAACTTATGATGTCTTTTCCCTGAGATGGGGGGCTCAGAGGGGGGAAGGCCTGTGGCCTGCATG  $\tt CTTCCCGATGGCCCACGGCAGGAGGTGTGTGGAAGTGTAAGGCCTAAGATGCTCACAGAGGTCCCTCATGACCTC$ CCTTCCCCAACTCCCGAATCCTCTTGAGTGTGGACCTCAACACCTTGAGCCCTAGTAAAGGAACTATGCAAAT GCAGGCCACTCTCCCCACCACGTCTGTGCCCCGCACTGTCCCCACAGCCTTCCACACCCTGTGCATAGGCAGCCC CTGCATTTGAATGATTTCCTTAGGATGAATGGTCCCTACACAAAGAGGCCTTGTGGGCCAAACCTGGAGAACCCTC CTAAATCCATAGAGTTTTCAAAATGTGAATCTTTGGAAGCCTTGAGTTCAGAATCTGCTGCTCTGGAATATTTCC AAAACAGGGAGTTGAAGTCCAACCTATTTAAAAACCCCACCATTTGGAGAATTACAAGGGTTTTGTCCTGAATTG TAGTGTTGGCAAGCCCAAGCCACTCGTGCTAACTGCTTTTTGTCTCGGTTGCTATTCCAAGAACAGAAGGAGGAA GTTGGCCAATTACAGCGTGTGTGCATGGATGTGTGTGGGGGGGCGTGCCTCTCAGAAACGCGGCCAGAAGACAAGC AGGGAAGTGAAAGGTCCCAGGCACACCCTGCCCATTGCAGGTGGCTCTTACAGCTCTCTGGTGCCAGCACGG ATCCCTGAAGTGACTCAGCCAGGCAGACATGAGACATGGCGGAGTGTCCAAATGGATCCTTTATTGGTGGTAGAG CAAAAAAACCCAAACACGATAAACCTTTCAAAAGACTTTCTAAGGATGATATTGGAATGCACCAGCCCTCACATG 

# 351/2825 FIGURE 323

 ${\tt MSGSFELSVQDLNDLLSDGSGCYSLPSQPCNEVTPRIYVGNASVAQDIPKLQKLGITHVLNAAEGRSFMHVNTNANFYKDSGITYLGIKANDTQEFNLSAYFERAADFIDQALAQKNGRVLVHCREGYSRSPTLVIAYLMMRQKMDVKSALSIVRQNREIGPNDGFLAQLCQLNDRLAKEGKLKP$ 

#### 352/2825 FIGURE 324

GAATTCGGGCGGGCTTCTTCGCTGCCGACGTACGACGAGTGGCCGGGCTCTTGCGTCTGGTAACGCGCTGTCTCT AACGCCAGCGCCGTCTCGCGCGCACTGCGCACAGACCACCCGCAGACGCCCGGCAGTCCGCAGGCCCAAACGCGC ACGCGACCCGCTCTCCGCACCGTACCCGGCCGCCTCGGC ATG GCGCCCCGCAGCGCCCCGGCGACCCCTGCTGCTGCTACTGCCTGTTGCTGCTCGGCCTCATGCATTGTCGTCAGCAGCCATGTTTATGGTGAAAAATGGCAACGG GACCGCGTGCATAATGGCCAACTTCTCTGCTGCCTTCTCAGTGAACTACGACACCAAGAGTGGCCCCAAGAACAT GACCTTTGACCTGCCATCAGATGCCACAGTGGTGCTCAACCGCAGCTCCTGTGGAAAAGAGAACACTTCTGACCC TCAGCTCATGAGTTTTGTTTATAACTTGTCAGACACACCCTTTTCCCCAATGCGAGCTCCAAAGAAATCAAGAC TGTGGAATCTATAACTGACATCAGGGCAGATATAGATAAAAAATACAGATGTGTTAGTGGCACCCAGGTCCACAT GAACAACGTGACCGTAACGCTCCATGATGCCACCATCCAGGCGTACCTTTCCAACAGCAGCTTCAGCAGGGGAGA GACACGCTGTGAACAAGACAGGCCTTCCCCAACCACGCGCCCCTGCGCCACCCAGCCCCTCGCCCTCACCCGT GCAGCTGAACCTCACCTATGAGAGGAAGGACAACACGACGGTGACAAGGCTTCTCAACATCAACCCCAACAAGAC  $\tt CTCGGCCAGCGGGAGCTGCGCCCCACCTGGTGACTCTGGAGCTGCACAGCGAGGGCACCACCGTCCTGCTCTT$  $\tt CCAGTTCGGGATGAATGCAAGTTCTAGCCGGTTTTTCCTACAAGGAATCCAGTTGAATACAATTCTTCCTGACGC$ CAGAGACCCTGCCTTTAAAGCTGCCAACGGCTCCCTGCGAGCGCTGCAGGCCACAGTCGGCAATTCCTACAAGTG CAACGCGGAGGAGCACGTCCGTGTCACGAAGGCGTTTTCAGTCAATATATTCAAAGTGTGGGTCCAGGCTTTCAA GGTGGAAGGTGGCCAGTTTGGCTCTGTGGAGGAGTGTCTGCTGGACGAGAACAGCACGCTGATCCCCATCGCTGT GGGTGGTGCCTGGCGGGGCTGGTCCTCATCGTCCTCATCGCCTACCTCGTCGGCAGGAAGAGAGGAGTCACGCAGG TCCTGTCGAAGGGGAGGCACACTTTCTGCAAACGTTTCTCAAATCTGCTTCATCCAATGTGAAGTTCATCTTGCA GCATTTACTATGCACAACAGAGTAACTATCGAAATGACGGTGTTAATTTTGCTAACTGGGTTAAATATTTTGCTA  $\tt CTTTCTCTGAGGGGGTGGGGGTGCCGCTCTCTCTGAGGGGGTGGGGGTGCTCTCTCCGAGGGGTGGAATGCC$ GCTGTCTCTGAGGGGTGGGGGTGCCGCTCTAAATTGGCTCCATATCATTGAGTTTAGGGTTCTGGTGTTTGGTTT GTCACTCGTGTGAGGCTGACATGCTCACACATTACAACAGTAGAGAGGGAAAATCCTAAGACAGAGGAACTCCAG AGTGGCGGCACGTCCTTGGCGTCTCTAATGTCTGCAGCTCAAGGGCTGGCACTTTTTTAAATATAAAAATGGTGT AATGTGAGATCGGTGCGTTCTCCTGATGTTTTGCCGTGGCTTGGGGGATTGTACACGGGACCAGCTCACGTAATGC ATTGCCTGTAACAATGTAATAAAAAGCCTCTTTCTTTCAAAAAAACCCCGAATTC

### 353/2825 FIGURE 325

MAPRSARRPLLLLLPVAAARPHALSSAAMFMVKNGNGTACIMANFSAAFSVNYDTKSGPKNMTFDLPSDATVVLN RSSCGKENTSDPSLVIAFGRGHTLTLNFTRNATRYSVQLMSFVYNLSDTHLFPNASSKEIKTVESITDIRADIDK KYRCVSGTQVHMNNVTVTLHDATIQAYLSNSSFSRGETRCEQDRPSPTTAPPAPPSPSPSPVPKSPSVDKYNVSG TNGTCLLASMGLQLNLTYERKDNTTVTRLLNINPNKTSASGSCGAHLVTLELHSEGTTVLLFQFGMNASSSRFFL QGIQLNTILPDARDPAFKAANGSLRALQATVGNSYKCNAEEHVRVTKAFSVNIFKVWVQAFKVEGGQFGSVEECL LDENSTLIPIAVGGALAGLVLIVLIAYLVGRKRSHAGYQTI

### 354/2825 FIGURE 326A

AAGCGGCTCGGGCTGCCTCAGAGTGCCGCGGGGGGCGTGGGGCGGTGCTGAGGAGCCGTGGCC GCAGCAGCAGCCCTCGCCGTTCGCGGAGCGCAGCCGAGCCGACCGTTGTCGATGCCGCTGAATGGGC TGAAGGAGGAGGACAAAGAGCCCCTCATCGAGCTCTTCGTCAAGGCTGGCAGTGATGGTGAAAGCATAGGAAACT GCCCCTTTTCCCAGAGGCTCTTCATGATTCTTTGGCTCAAAGGAGTTGTATTTAGTGTGACGACTGTTGACCTGA AAACGGATGTAAATAAGATTGAGGAATTTCTTGAAGAGGTCTTATGCCCTCCCAAGTACTTAAAGCTTTCACCAA AACACCCAGAATCAAATACTGCTGGAATGGACATCTTTGCCAAATTCTCTGCATATATCAAGAATTCAAGGCCAG AGGCTAATGAAGCACTGGAGAGGGGTCTCCTGAAAACCCTGCAGAAACTGGATGAATATCTGAATTCTCCTCTCC CTGATGAAATTGATGAAAATAGTATGGAGGACATAAAGTTTTCTACACGTAAATTTCTGGATGGCAATGAAATGA CATTAGCTGATTGCAACCTGCTGCCCAAACTGCATATTGTCAAGGTGGTGGCCAAAAAATATCGCAACTTTGATA CCAGTGATAAGGAGGTTGAAATAGCATATAGTGATGTAGCCAAAAGACTCACCAAG**TAA**AATCGCGTTTGTAAAA TTTTGCACGAACATGCAGTTATTGAAGATTAGGATCAAGGATAGACAAGGTATAGTAGTTATCTTAAAATATACA CTCCTAAGCAGTATTATTTTAAAATCCTTTACCCTGGCTACCTCCCCTACCCGGGTTCCCCTCTCTTTAATTTGG AGACACTCCACCACAAACTTTTCACTTTAGAGGTAGCTTGCCATCTCTCAGGAGCCCTCACCATTGTGTCCATTC ACTGTGTATAGATGGCAGAACTTTTGAGGTGCAATGTTTAATTGTTAAAAATAGTAGCCACGACTTTATCAGGCA GTATATGAAAGGATTGCTACTGTATCAGAAATATTGTTTCAATTTAGTCTATCCTGGATATGTACTAACGAATAT TTTGCCTAAAAGCATTTATCCTTCATACCAATTGTAACATCTGACACCATGTAGAAGCTAAAAGTTTAGAGGGAG TGAGCGTTTTCTCAAGACCTTCCTCAAGCATTTTATCTTTAGAAGAGAAACTGATGGGCACCTGATACTCTGTCT TGTTTTTACGTAATCATAGAGCTTGCTATTTGTACATCTGTTGAGCAACACTACATAACTGATTTTTAGTTGACT TATAAAAGAAAGATAAAACACAGGTCACCAATTTTCTCATTTCACCCCATTTACCTTGTATAGAGGATTGTTCAT TTTAATTGAAGCCCTAAACCAGGAATTATTTGTGTTCTAACAGGAGGATGAACTTGCTGAAAATAAAACTTTGCT ATGTATTTACTCTTTTTTAAAAGACAAAAGCAAAACCAGACTTTCTACGTACTACTCCAAAGACTGTGATTGTGA CTATAATACATTTTTGGTAATTTTTTTATACCTAATTTGTATAGGAAGTGCTATTTCTCATAGGCTGTTTCTTGA AATTTTAAGTTTATTGCTTTAAAATGGCAGTGTTTCTCCCACTTTGATATGCTAACATTTAGTAAGCACTGGCTT AATCTCCATGTTGTATCCATTTGGCTCAGGAAGATTCTTTGCCTTACCTTTCTTAGAACTCTTTATTGCTTATCA AAAGTTTGAGTACCCGCTTGGTTTTTTTTTGGTAATTAAATATTGTATGATTTATCTGGTTCAAGGAAGATGCAC TATTCAGTTATCTATTGAGAAATTATTTTGCAGTGGTTTTAGTGGGTGAAAATGTCCCATCTGCACCAGTACACA CCTCATAGCTTAAAGTATAAAAAAAGATTCAAGAGCAGTGAGGTTTGTTCTTTCCAGTGAATGGTGGACTGAGTG GTGCGAGGTGGAGGGCTAACAAGAGGAAAGAACTACATTCTTCAGAATACAGTGATGAAAATTCATTTTGAAACT CAAATATTTTCATTTTGGATATTCTCCTGTTTTTATTAAACCAGTGATTACACCTGGCCATCCCTCTAAATGTTC TAGGAAGGCATGTCTATTGTGATTTTGATGAAGACAGAATTATTTTTCTCTGTAGAAACACAGATACCACTTTAT ATGCTTTAACCCCATTGTATGTTTTGTGGAAAGAGCATAGTTTAACATCTTGAGAAATTTGGGACATAAAGTTTTC ATGGTAGACAGTTCATGCAGTATATGAATTGACATAATGGAAATAATCTGATTTTATTTTTACAACTAACATCCA TTCCCCTTCATTTAAACACCTTTTGTGTTTTACTTCAGTGAGGAGATTGGAGTCTGAATGGATCTGTTTTCCAAG AGATTCTGAGAAATTTTTGTATTCAGCAGTTGGAAAGCTCTCTATTCTAGTTGATAAAACTTCCCTTTTTTGATG

# 355/2825 FIGURE 326B

# 356/2825 FIGURE 327

MALSMPLNGLKEEDKEPLIELFVKAGSDGESIGNCPFSQRLFMILWLKGVVFSVTTVDLKRKPADLQNLAPGTHP PFITFNSEVKTDVNKIEEFLEEVLCPPKYLKLSPKHPESNTAGMDIFAKFSAYIKNSRPEANEALERGLLKTLQK LDEYLNSPLPDEIDENSMEDIKFSTRKFLDGNEMTLADCNLLPKLHIVKVVAKKYRNFDIPKEMTGIWRYLTNAY SRDEFTNTCPSDKEVEIAYSDVAKRLTK

## 357/2825 FIGURE 328

TGCTGCGAACCACGTGGGTCCCGGGCGCGTTTCGGGTGCTGGCGGCTGCAGCCGGAGTTCAAACCTAAGCAGCTG
GAAGGAACCATGGCCAACTGTGAGCGTACCTTCATTGCGATCAAACCAGATGGGGTCCAGCGGGGTCTTGTGGGA
GAGATTATCAAGCGTTTTGAGCAGAAAGGATTCCGCCTTGTTGGTCTGAAATTCATGCAAGCTTCCGAAGATCTT
CTCAAGGAACACTACGTTGACCTGAAGGACCGTCCATTCTTTGCCGGCCTGGTGAAATACATGCACTCAGGGCCG
GTAGTTGCCATGGTCTGGGAGGGCCTGAATGTGGTGAAGACGGGCCGAGTCATGCTCGGGGAGACCAACCCTGCA
GACTCCAAGCCTGGGACCATCCGTGGAGACTTCTGCATACAAGTTGGCAGGAACATTATACATGGCAGTGATTCT
GTGGAGAGTGCAGAAAGGAGATCGGCTTGTGGTTTCACCCTGAGGAACTGGTAGATTACACGAGCTGTGCTCAG
AACTGGATCTATGAATGAATGACAGGAGGCAGACCACCATTGCTTTTCACATCCATTTCCCCTCCTTCCCATGGGCAGA
GGACCAGGCTGTAGGAAATCTAGTTATTTACAGGAACTTCATCATAATTTGGAGGGAAGCTCTTGGAGCTGTGAG
TTCTCCCTGTACAGTGTTACCATCCCGACCATCTGATTAAAATGCTTCCTCCCAGC

# 358/2825 FIGURE 329

MANCERTFIAIKPDGVQRGLVGEIIKRFEQKGFRLVGLKFMQASEDLLKEHYVDLKDRPFFAGLVKYMHSGPVVA MVWEGLNVVKTGRVMLGETNPADSKPGTIRGDFCIQVGRNIIHGSDSVESAEKEIGLWFHPEELVDYTSCAQNWI YE

### 359/2825 FIGURE 330A

GCGGCGGTGGCGGGAGACCCGAAC<u>ATG</u>GCGACCGCGCGCACCTTCGGGCCCGAGCGGGAAGCCGAGCCGGCCA AGGAAGCGCGCGTCGTGGGCTCGGAGCTTGTGGACACTTATACGGTTTACATCATCCAGGTCACTGATGGCAGCC ATGAGTGGACAGTAAAGCACCGCTACAGCGACTTCCATGACCTGCATGAAAAGCTCGTTGCAGAGAAAAGATTG ATAAAAACCTGCTTCCGCCCAAAAAGATAATTGGGAAAAACTCAAGAAGCTTGGTGGAGAAGAGGGAGAAGGATC TGGAGGTCTACCTCCAGAAGCTCCTGGCTGCCTTCCCTGGCGTGACCCCCAGAGTACTGGCCCACTTCTTGCATT TTCACTTCTATGAGATAAATGGCATCACCGCGGCACTGGCTGAAGAGCTCTTTGAGAAAGGAGAACAGCTCCTGG GGGCCGGCGAGGTCTTTGCCATTGGACCCCTGCAGCTGTATGCCGTCACGGAGCAGCTGCAGCAGGGAAAGCCCA CGTGCGCCAGTGGGGATGCCAAGACCGACCTCGGGCACATCCTGGACTTCACCTGTCGCCTTAAGTACCTTAAGG TTTCTGGCACAGAAGGACCTTTTGGGACCAGCAACATTCAGGAGCAGCTCCTGCCGTTCGACCTATCAATATTCA AGTCCCTGCATCAGGTGGAGATAAGTCACTGTGATGCTAAGCACATCAGAGGGCTGGTCGCATCGAAGCCCACCT TTGACCTGAGCCACACAGCATCTCCGAGATCGACGAGTCTGTGAAACTGATCCCAAAGATTGAGTTCCTGGACC TGAGTCACAATGGATTGCTGGTTGTGGACAATCTGCAGCACCTGTATAACCTTGTGCATCTGGACCTGTCCTACA TAGAGAGTCTGAGTGGCCTGCACAAGCTCTACTCACTGGTCAACCTGGATCTCCGGGACAACAGGATCGAACAGA TGGAGGAGGTCCGGAGCATAGGCAGCCTCCCGTGTCTGGAGCACGTGTCTCTGCTGAACAACCCTCTGAGCATCA CCACAGAGAGGAGCTGGACACTGTGGAAGTGCTGAAAGCAATTCAGAAAGCCAAGGAGGTCAAGTCCAAACTGA GCAACCCAGAGAAGAAGGGTGGTGAAGACTCCCGGCTCTCAGCTGCCCCCTGCATCAGACCCAGCAGCTCCCCTC AGGCCCTGGCCAGCAGCCTCTCGTCCACTGACAGTCTGACTCCCGAGCACCAGCCCATTGCCCAGGGATGTTCTG ATTCCTTGGAGTCCATCCCTGCGGGACAGCCTCCCGATGATTTAAGGGACGTGCCAGGAGCTGTTGGTGGTG  $\tt CAAGCCCAGAACATGCCGAGCCGGAGGTCCAGGTGGTGCCGGGGTCTGGCCAGATCATCTTCCTGCCCTTCACCT$ GCATTGGCTACACGGCCACCAATCAGGACTTCATCCAGCGCCTGAGCACACTGATCCGGCAGGCCATCGAGCGGC AGCTGCCTGCATCGAGGCTGCCAACCAGCGGGAGGAGGGCCAGGGTGAACAGGGCGAGGAGGAGGATGAGG GAGGAGGCCAGGGGGAGGAAGAGGAGGAAGAGGAGGATGAAGAGGCCGAGGAGGAGCGCCTGGCTCTGGAAT ATGTGCAGGCCAGTATCCGCCAGTTCGCCGCCTTGTGCTCACCGACTTCGGCATCGCAGTCTTCGAGATCC CGCACCAGGAGTCTCGGGGCAGCAGCCAGCACATCCTCTCCCTGCGCTTTGTCTTTTGCTTCCCGCATGGCG ACCTCACCGAGTTTGGCTTCCTCATGCCGGAGCTGTGTCTGGTGCTCAAGGTACGGCACAGTGAGAACACGCTCT TCATTATCTCGGACGCCCCAACCTGCACGAGTTCCACGCGGACCTGCGCTCATGCTTTGCACCCCAGCACATGG  ${\tt CCATGCTGTGTAGCCCCATCCTCTACGGCAGCCACACCAGCCTGCAGGAGTTCCTGCGCCAGCTGCTCACCTTCT}$ ACAAGGTGGCTGGCGGCTGCCAGGAGCGCAGCCAGGGCTGCTTCCCCGTCTACCTGGTCTACAGTGACAAGCGCA TGGTGCAGACGGCCGCGGGGACTACTCAGGCAACATCGAGTGGGCCAGCTGCACACTCTGTTCAGCCGTGCGGC GCTCCTGCTGCGCCCCTCTGAGGCCGTCAAGTCCGCCGCCATCCCCTACTGGCTGTTGCTCACGCCCCAGCACC TCAACGTCATCAAGGCCGACTTCAACCCCATGCCCAACCGTGGCACCACAACTGTCGCAACCGCAACAGCTTCA AGCTCAGCCGTGTGCCGCTCTCCACCGTGCTGCTGCACCCCACACGCAGCTGTACCCAGCCTCGGGGCGCCTTTG CTGATGGCCACGTGCTAGAGCTGCTCGTGGGGTACCGCTTTGTCACTGCCATCTTCGTGCTGCCCCACGAGAAGT TCCACTTCCTGCGCGTCTACAACCAGCTGCGGGCCTCGCTGCAGGACCTGAAGACTGTGGTCATCGCCAAGACCC CCGGGACGGGAGCCCCCAGGGCTCCTTTGCGGATGGCCAGCCTGCCGAGCGCAGGGCCAGCAATGACCAGC GTCCCAGGAGGTCCCAGCAGAGGCTCTGGCCCCGGCCCCAGTGGAAGTCCCAGCTCCAGCCCCTGCAGCAGCCCT CAGCCTCAGGCCCAGCGAAGACTCCGGCCCCAGCAGAGGCCTCAACTTCAGCTTTGGTCCCAGAGGAGACGCCAG TGGAAGCTCCAGCCCACCCCAGCCGAGGCCCCTGCCCAGTACCCGAGTGAGCACCTCATCCAGGCCACCTCGG TCATCGAGCTCTTCCACAGCAGCATTGCTGAGGTTGAAAACGAGGAGCTGAGGCACCTCATGTGGTCCTCGGTGG TGTTCTACCAGACCCCAGGGCTGGAGGTGACTGCCTGCGTGCTCCTCCACCAAGGCTGTGTACTTTGTGCTCC

### 360/2825 FIGURE 330B

ACGACGCCTCCGCCGCTACTTCTCAGAGCCACTGCAGGATTTCTGGCATCAGAAAAACACCGACTACAACAACA GCCCTTTCCACATCTCCCAGTGCTTCGTGCTAAAGCTTAGTGACCTGCAGTCAATGTGGGGGCTTTTCGACC AGCATTTCCGGCTGACGGGTTCCACCCCGATGCAGGTGGTCACGTGCTTGACGCGGGACAGCTACCTGACGCACT TCTACTCCGAGTTTGGGAACAAGACCACAGGGAAGATGGAGAACTACGAGCTGATCCACTCTAGTCGCGTCAAGT TTACCTACCCCAGTGAGGAGGAGATTGGGGACCTGACGTTCACTGTGGCCCAAAAGATGGCTGAGCCAGAGAAGG  $\verb|CCCAGCCCTCAGCATCCTGCTGTACGTGCAGGCCTTCCAGGTGGCCATGCCACCCCTGGGTGCTGCAGGGGCC| \\$ TGCCCGAGTTTGCCAAAGAGCCGCCGCAGAGAGACAGGTACCGGCTGGACGATGGCCGCCGCGTCCGGGACCTGG ACCGAGTGCTCATGGGCTACCAGACCTACCCGCAGGCCCTCACCCTCGTCTTCGATGACGTGCAAGGTCATGACC  $\verb|ACGCCTACTGGGGCAGGCCAGGCTTTTGTGTTCTCTAAAAATGTTTTATCCTCCCTTTGGTACCTTAATTTG|$ GCCCTCAGGGCTGTCGGTGTCTCAGCCTCCCACAGGTGGTACAGCCGTGCACACCAGTGTCGTGTCTGCTG TTGTGGGACCGTTGTTAACACGTGACACTGTGGGTCTGACTTTCTCTTCTACACGTCCTTTCCTGAAGTGTCGAG CAGAATGCACATTGGAAGCTCCCACCCCATATTGTTCTTCAAAGTGGAGGTCTCCCCTGATCCAGACAAGTGGGA GAGCCCGTGGGGGCAGGGGACCTGGAGCTGCCAGCACCAAGCGTGATTCCTGCTGCCTGTATTCTATTCCAAT AAAGCAGAGTTTGACACCGTCTTAAAAAAAAA

### 361/2825 FIGURE 331

MATARTFGPEREAEPAKEARVVGSELVDTYTVYIIQVTDGSHEWTVKHRYSDFHDLHEKLVAERKIDKNLLPPKK IIGKNSRSLVEKREKDLEVYLQKLLAAFPGVTPRVLAHFLHFHFYEINGITAALAEELFEKGEQLLGAGEVFAIG PLQLYAVTEQLQQGKPTCASGDAKTDLGHILDFTCRLKYLKVSGTEGPFGTSNIQEQLLPFDLSIFKSLHQVEIS HCDAKHIRGLVASKPTLATLSVRFSATSMKEVLVPEASEFDEWEPEGTTLEGPVTAVIPTWQALTTLDLSHNSIS EIDESVKLIPKIEFLDLSHNGLLVVDNLOHLYNLVHLDLSYNKLSSLEGLHTKLGNIKTLNLAGNLLESLSGLHK LYSLVNLDLRDNRIEQMEEVRSIGSLPCLEHVSLLNNPLSIIPDYRTKVLAQFGERASEVCLDDTVTTEKELDTV EVLKAIQKAKEVKSKLSNPEKKGGEDSRLSAAPCIRPSSSPPTVAPASASLPQPILSNQGIMFVQEEALASSLSS TDSLTPEHQPIAQGCSDSLESIPAGQAASDDLRDVPGAVGGASPEHAEPEVQVVPGSGQIIFLPFTCIGYTATNQ DFIQRLSTLIRQAIERQLPAWIEAANQREEGQGEGEEEDEEEEEEDVAENRYFEMGPPDVEEEEGGGQGEEEE EEEEDEEAEEERLALEWALGADEDFLLEHIRILKVLWCFLIHVOGSIROFAACLVLTDFGIAVFEIPHOESRGSS QHILSSLRFVFCFPHGDLTEFGFLMPELCLVLKVRHSENTLFIISDAANLHEFHADLRSCFAPQHMAMLCSPILY GSHTSLQEFLRQLLTFYKVAGGCQERSQGCFPVYLVYSDKRMVQTAAGDYSGNIEWASCTLCSAVRRSCCAPSEA VKSAAIPYWLLLTPOHLNVIKADFNPMPNRGTHNCRNRNSFKLSRVPLSTVLLDPTRSCTOPRGAFADGHVLELL VGYRFVTAIFVLPHEKFHFLRVYNQLRASLQDLKTVVIAKTPGTGGSPQGSFADGQPAERRASNDQRPQEVPAEA LAPAPVEVPAPAPAAASASGPAKTPAPAEASTSALVPEETPVEAPAPPPAEAPAQYPSEHLIQATSEENQIPSHL PACPSLRHVASLRGSAIIELFHSSIAEVENEELRHLMWSSVVFYQTPGLEVTACVLLSTKAVYFVLHDGLRRYFS EPLQDFWHQKNTDYNNSPFHISQCFVLKLSDLQSVNVGLFDQHFRLTGSTPMQVVTCLTRDSYLTHCFLQHLMVV LSSLERTPSPEPVDKDFYSEFGNKTTGKMENYELIHSSRVKFTYPSEEEIGDLTFTVAOKMAEPEKAPALSILLY VQAFQVGMPPPGCCRGPLRPKTLLLTSSEIFLLDEDCVHYPLPEFAKEPPQRDRYRLDDGRRVRDLDRVLMGYQT YPQALTLVFDDVQGHDLMGSVTLDHFGEVPGGPARASQGREVQWQVFVPSAESREKLISLLARQWEALCGRELPV ELTG

# 362/2825 FIGURE 332

CCCGGAGGTAGGCGGGCGCGGGCCTGTTGGGTCTTTGGGACGCGGGTCCCGCTGGGGCCGGGGATGCTCCTCTC CCCTTCTTCCAGGAAGCGCCTCTTGGACGCGTGTGACCGATGCCCAGATTGCACGACCACTTCTGGAGCTGCTCCTGTGCGCACAGCGCGAGGCCCCCGCGCGAGCCAGCACCGCGGGGCTGCCGCCCAAGGTTGGGGAGATG ATCAACGTTTCCGTGTCCGGGCCCTCCCTGCTGGCGCCCACGGTGCCCCGGACGCTGACCCCGCGCCCAGGGGC CGCAGTGCTGCGATGAGCGGCCCCGAGCCCGGCAGCCCCTACCCCAACACCTGGCATCATCGCCTGTTGCAGAGG AGCCTCGTGCTCTTCTCGGTTGGGGTGGTCCTAACCCTGGTGCTCAACCTGCTGCAGATCCAGAGGAATGTCACT GCTGTTGTTGGCCTACTGTACCCCTGTATCGACAGTCACCTCGGAGAACCCCACAAATTTAAGAGAGAATGGGCC AGTGTCATGCGCTGCATAGCAGGTTTTGGTGGCATTAACCACGCCAGTGCTAAATTGGATTTTGCCAATAATGTC CAGCTGTCCTTGACTTTAGCAGCCCTATCTTTGGGCCTTTGGTGGACATTTGATCGTTCCAGAAGTGGCCTTGGG CTGGGGATCACCATAGCTTTTCTAGCTACGCTGATCACGCAGTTTCTCGTGTATAATGGTGTCTATCAGTATACG TCCCCAGATTTCCTCTATATTCGTTCTTGGCTCCCTTGTATATTTTTCTCAGGAGGCGTCACGGTGGGGAACATA GGACGACAGTTAGCTATGGGTGTTCCTGAAAAGCCCCATAGTGAT**TGA**GTCTTCAAAACCACCGATTCTGAGAGC TCGGGCTGACTGTACAAATGACTCCTGGAAAAAACTCGTCACCTATCTAGAAAAGTCAAGAATAGGGAGGTGGAG GCTTAACAAACATGAAGTATGGTGTAATAGGAATAATATTTATCCAAAAGATTTTTAAAAAATAGGGCTGTGTTTA AAGAAGGAATCAAAACAAGAAAAGCAGCAGTGATTATAGAGAGGTCACACTCTAAGTGGGGTCGCGGCGTGGCCA AACTGCTTGTCATTCCTCCCGCGGCAGTGTTGTGTAGACAATCTTACTGAGCAAAAGGCAATGAAAAGTCTTGGG CTCCCACACTGCGATATATTGGAATTCCCACCTCAGTTTATGAAGTTTATTTCGAAATCCATAGTCATCTAAGAA TGTTTTTTGGCACAGCCCTATGGAATTTGCAATCTGTGATTGCCTTGTAAAAAGGAGAGTGCATATGGCACTGCA TTAAACGTGTGGTGTTTCTAGTCAATGATATTGGTGAGCACAATGTATTCATTTAATGGCATAGACCATACCAGA CCTAATTTGCAAGTATTGGGTCTTCAAACTTCAAGTGCAATGTATTATGAAAACCAATCTGAGCCTTGTATCTCT AAACTCGGCAATGATTTCTTCAGTTGTGAAGTTCCTTTCGGGTTACAACCTCCACTGGAACCCTCAACCTTCGAA ATACTCCAGTTTTGGGGGTTGGGGCCATTTACTTATAAATTTACCGCCGGGTTTTTGGAATCTACATGTCTTGGG GGCGGGCTCAAATTCTTCGAAAGTGGTTGGATTAAAA

# 363/2825 FIGURE 333

MPRLHDHFWSCSCAHSARRRGPPRASTAGLPPKVGEMINVSVSGPSLLAAHGAPDADPAPRGRSAAMSGPEPGSP YPNTWHHRLLQRSLVLFSVGVVLTLVLNLLQIQRNVTLFPEEVIATIFSSAWWVPPCCGTAAAVVGLLYPCIDSH LGEPHKFKREWASVMRCIAGFGGINHASAKLDFANNVQLSLTLAALSLGLWWTFDRSRSGLGLGITIAFLATLIT QFLVYNGVYQYTSPDFLYIRSWLPCIFFSGGVTVGNIGRQLAMGVPEKPHSD

### 364/2825 FIGURE 334A

GCTGCCCGCCGAAGCGAAACCGAGTCCGGGCCCGTCCCTCCGCGGCCCCATCCGCCCGGTGCACCCGGGG GTGACCAGCGTCACGCTGCTGGTGGCCCGGGTCTTCCAGAAGGCTGTAGACCAGAGTATCGAGAAGAAAATTGTG TTAAGGAATGGTACTGAGGCATTTGACTCCTGGGAGAAGCCCCCTCTGCCTGTGTATACTCAGTTCTATTTCTTC AATGTCACCAATCCAGAGGAGATCCTCAGAGGGGAGACCCCTCGGGTGGAAGAAGTGGGGCCATACACCTACAGG GAACTCAGAAACAAAGCAAATATTCAATTTGGAGATAATGGAACAACAATATCTGCTGTTAGCAACAAGGCCTAT GTTTTTGAACGAGACCAATCTGTTGGAGACCCTAAAATTGACTTAATTAGAACATTAAATATTCCTGTATTGACT GTCATAGAGTGGTCCCAGGTGCACTTCCTCAGGGAGATCATCGAGGCCATGTTGAAAGCCTATCAGCAGAAGCTC TTTGTGACTCACAGTTGACGAATTGCTCTGGGGCTACAAAGATGAAATCTTGTCCCTTATCCATGTTTTCAGG CCCGATATCTCTCCCTATTTTGGCCTATTCTATGAGAAAAATGGGACTAATGATGAGAGACTATGTTTTTCTAACT GGAGAAGACAGTTACCTTAACTTTACAAAAATTGTGGAATGGAATGGGAAAACGTCACTTGACTGGTGGATAACA GACAAGTGCAATATGATTAATGGAACAGATGGAGATTCTTTTCACCCACTAATAACCAAAGATGAGGTCCTTTAT TTTCGGTATAAAGTTCCTGCAGAAATATTAGCCAATACGTCAGACAATGCCGGCTTCTGTATACCTGAGGGAAAC TGCCTGGGCTCAGGAGTTCTGAATGTCAGCATCTGCAAGAATGGTGCACCCATCATTATGTCTTTCCCACACTTT TACCAAGCAGATGAGAGGTTTGTTTCTGCCATAGAAGGCATGCACCCAAATCAGGAAGACCATGAGACATTTGTG GACATTAATCCTTTGACTGGAATAATCCTAAAAGCAGCCAAGAGGTTCCAAATCAACATTTATGTCAAAAAATTA GATGACTTTGTTGAAACGGGAGACATTAGAACCATGGTTTTCCCAGTGATGTACCTCAATGAGAGTGTTCACATT GATAAAGAGACGGCGAGTCGACTGAAGTCTATGATTAACACTACTTTGATCATCACCAACATACCCTACATCATC ACAGCGGATGAAAGAGCACCCCTCATTCGAACC**TAA**ACATTGCCTTTGCTTGGTGAAGAAACTGTGTGAGCTGTC CTGACCTGGACGATGACGTGGGGAAACCCTCCACCTCCTTGCAGGCTTGTTGCCTGTTGAAAGAAGAAAAAAGAC ATAAAATCATGTGGGCTCTGAAATTGTTCTTTTATGTGTCTAGCAAGTATTTAATAAACCCTTGTATAGTAATTT TGTTGTTGTTGGGTGCTGGTAGCTCCAGAATTTTGTGACCACTATTGTGGGTAAAATGTCTCTGCATCACTTGTT AATGCTACTGGTCTAACTTCATTCAGTATGCTTCATTCACCGAACTTTGTGCTCAAAATGCGTATATACCATTTT ATGTTGTATTCCTCCATTTCACTTGCAAAACAGAAGTAAATAAGAGTTCGGGACCCAGGGTAAAATGGTAGCTTC ATCCAATATATCATTCAAATGCATCTGATTTCTAAAACATATTACATTTTATGCTGATCTTCAGTTCATAATTCT TACCCATGAGTTGATAGTGATAAGCTCCTAAGGATTTTTAACTTGTACTTTTTGTGAACGAAGAGAATGCATAAAT GTTTTTTTTTTTTGTCAAAATGATCATGCTACGAAAAGATGCTTCTGAGAGAATGTAATGAGTAACTGATTTTTCT TATGGGAAATCCATCTATCTACAGCCTAAGTTACACATAAGTTTCAGAAAGTCTGATTAGACTAAAGAGATATTT GTGTTTTGTAGCTCACTGTCACTCACTGAATAGAGAAACGTGTGCCCTATACTTCCTGTGACAATCATTTTGCTG CACTTTTCAAAGGACTTCTTGGTTTCAGCATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTC GATTTCAATGTAAGCAAATGAAGAAAAAATATAGATTTCATTTCCATAGCTTCTTATCCCTGTATTGAGGTAAT AAATTGTTTTACTGACAATTTTTCCTTTTTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCA ATTCATTAAACTCTCAAATTTTCTATAGAATCAAGATAGAACCTTTAGATACTCCAACTCACCAAAATGTAAAAA AACTAACAAAATATTTGGTCTTCAATAATGCTAAATATCTACATTTTTAGAATTTATCAACATTTAACTAGATA ATTGGGCATGTCTTAATTATGCATGTACTTATCCATACTAATAAAATTGACAATGCTAGTGCATACTTATTGGTT

# 365/2825 FIGURE 334B

# 366/2825 FIGURE 335

MGRCCFYTAGTLSLILLLVTSVTLLVARVFQKAVDQSIEKKIVLRNGTEAFDSWEKPPLPVYTQFYFFNVTNPEEI LRGETPRVEEVGPYTYRELRNKANIQFGDNGTTISAVSNKAYVFERDQSVGDPKIDLIRTLNIPVLTVIEWSQVH FLREIIEAMLKAYQQKLFVTHTVDELLWGYKDEILSLIHVFRPDISPYFGLFYEKNGTNDGDYVFLTGEDSYLNF TKIVEWNGKTSLDWWITDKCNMINGTDGDSFHPLITKDEVLYVFPSDFCRSVYITFSDYESVQGLPAFRYKVPAE ILANTSDNAGFCIPEGNCLGSGVLNVSICKNGAPIIMSFPHFYQADERFVSAIEGMHPNQEDHETFVDINPLTGI ILKAAKRFQINIYVKKLDDFVETGDIRTMVFPVMYLNESVHIDKETASRLKSMINTTLIITNIPYIIMALGVFFG LVFTWLACKGQGSMDEGTADERAPLIRT

## 367/2825 FIGURE 336A

GCGCGACCGTCCCGGGGGTGGGGCCGGGCGCGCGGCGAGGGGGGGAGGTGGCTGCGGTAGCAGCGCGCGG ACACTCGGGAGGACAACGTGATCCGGAAATATGGAGACCCCGGGAGCCTCTTCGGCTTCTCGCTGGCCATGCACT GGCAACTGCAGCCCGAGGACAAGCGGCTGTTGCTCGTGGGGGCCCCGCGCGGAGAAGCGCTTCCACTGCAGAGAG CCAACAGAACGGGAGGCTGTACAGCTGCGACATCACCGCCCGGGGGCCATGCACGCGGATCGAGTTTGATAACG ATGCTGACCCCACGTCAGAAAGCAAGGAAGATCAGTGGATGGGGGTCACCGTCCAGAGCCAAGGTCCAGGGGGCA AGGTCGTGACATGTGCTCACCGATATGAAAAAAGGCAGCATGTTAATACGAAGCAGGAATCCCGAGACATCTTTG GGCGATTGAGAGGCCATGAGAAATTTGGCTCTTGCCAGCAAGGTGTAGCAGCTACTTTTACTAAAGACTTTCATT ACATTGTATTTGGAGCCCCGGGTACTTATAACTGGAAAGGGATTGTTCGTGTAGAGCAAAAGAATAACACTTTTT TTGACATGAACATCTTTGAAGATGGGCCTTATGAAGTTGGTGGAGAGACTGAGCATGAAGATCTCGTTCCTG TTCCTGCTAACAGTTACTTAGGTTTTTCTTTGGACTCAGGGAAAGGTATTGTTTCTAAAGATGAGATCACTTTTG TATCTGGTGCTCCCAGAGCCAATCACAGTGGAGCCGTGGTTTTGCTGAAGAGAGACATGAAGTCTGCACATCTCC TCCCTGAGCACATATTCGATGGAGAAGGTCTGGCCTCTTCATTTGGCTATGATGTGGCGGTGGTGGACCTCAACA AGGATGGGTGGCAAGATATAGTTATTGGAGCCCCACAGTATTTTGATAGAGATGGAGAAGTTGGAGGTGCAGTGT ATGTCTACATGAACCAGCAAGGCAGATGGAATAATGTGAAGCCAATTCGTCTTAATGGAACCAAAGATTCTATGT TTGGCATTGCAGTAAAAAATATTGGAGATATTAATCAAGATGGCTACCCAGATATTGCAGTTGGAGCTCCGTATG GTATATCACCTTATTTTGGATATTCAATTGCTGGAAACATGGACCTTGATCGAAATTCCTACCCTGATGTTGCTG TTGGTTCCCTCTCAGATTCAGTAACTATTTTCAGATCCCGGCCTGTGATTAATATTCAGAAAACCATCACAGTAA CTCCTAACAGAATTGACCTCCGCCAGAAAACAGCGTGTGGGGCGCCTAGTGGGATATGCCTCCAGGTTAAATCCT GTTTTGAATATACTGCTAACCCCGCTGGTTATAATCCTTCAATATCAATTGTGGGCACACTTGAAGCTGAAAAAG AAAGAAGAAATCTGGGCTATCCTCAAGAGTTCAGTTTCGAAACCAAGGTTCTGAGCCCAAATATACTCAAGAAC TAACTCTGAAGAGGCAGAAACAGAAAGTGTGCATGGAGGAAACCCTGTGGCTACAGGATAATATCAGAGATAAAC TGCGTCCCATTCCCATAACTGCCTCAGTGGAGATCCAAGAGCCAAGCTCTCGTAGGCGAGTGAATTCACTTCCAG AAGTTCTTCCAATTCTGAATTCAGATGAACCCAAGACAGCTCATATTGATGTTCACTTCTTAAAAGAGGGATGTG GAGACGACAATGTATGTAACAGCAACCTTAAACTAGAATATAAATTTTGCACCCGAGAAGGAAATCAAGACAAAT TTTCTTATTTACCAATTCAAAAAGGTGTACCAGAACTAGTTCTAAAAGATCAGAAGGATATTGCTTTAGAAATAA CAGTGACAAACAGCCCTTCCAACCCAAGGAATCCCACAAAAGATGGCGATGACGCCCATGAGGCTAAACTGATTG CAACGTTTCCAGACACTTTAACCTATTCTGCATATAGAGAACTGAGGGCTTTCCCTGAGAAACAGTTGAGTTGTG TTGCCAACCAGAATGGCTCGCAAGCTGACTGTGAGCTCGGAAATCCTTTTAAAAGAAATTCAAATGTCACTTTTT ATTTGGTTTTAAGTACAACTGAAGTCACCTTTGACACCCCATATCTGGATATTAATCTGAAGTTAGAAACAACAA GCAATCAAGATAATTTGGCTCCAATTACAGCTAAAGCAAAAGTGGTTATTGAACTGCTTTTATCGGTCTCGGGAG  $\verb|TTGCTAAACCTTCCCAGGTGTATTTTGGAGGTACAGTTGTTGGCGAGCAAGCTAT | \dot{\texttt{G}}\texttt{AAATCTGAAGATGAAGTGG}$ GAAGTTTAATAGAGTATGAATTCAGGGTAATAAACTTAGGTAAACCTCTTACAAACCTCGGCACAGCAACCTTGA ACATTCAGTGGCCAAAAGAAATTAGCAATGGGAAATGGTTGCTTTATTTGGTGAAAGTAGAATCCAAAGGATTGG ACTGTAGCGTGAACGTGAACTGTGTGAACATCAGATGCCCGCTGCGGGGGCTGGACAGCAAGGCGTCTCTTATTT TGCGCTCGAGGTTATGGAACAGCACATTTCTAGAGGAATATTCCAAACTGAACTACTTGGACATTCTCATGCGAG CCTTCATTGATGTGACTGCTGCCGAAAATATCAGGCTGCCAAATGCAGGCACTCAGGTTCGAGTGACTGTGT TTCCCTCAAAGACTGTAGCTCAGTATTCGGGAGTACCTTGGTGGATCATCCTAGTGGCTATTCTCGCTGGGATCT  $\tt CCACATATCACAAGGCTGAGATCCATGCTCAGCCATCTGATAAAGAGAGGGCTTACTTCTGATGCA{\bf TAGT}$ CTACTTCTGTAATTGTGTGGATTCTTTAAACGCTCTAGGTACGATGACAGTGTTCCCCGATACCATGCTGTAAGG 

## 368/2825 FIGURE 336B

CAACTCAGAAATTCAATTTGGATTTAAAAGCCTGCTCAATCCCTGAGGACTGATTTCAGAGTGACTACACACAGT ACGAACCTACAGTTTTAACTGTGGATATTGTTACGTAGCCTAAGGCTCCTGTTTTGCACAGCCAAATTTAAAACT GTTGGAATGGATTTTTCTTTAACTGCCGTAATTTAACTTCTGGGTTGCCTTTGTTTTTTGGCGTGGCTGACTTAC ATCATGTGTTGGGGAAGGGCCTGCCCAGTTGCACTCAGGTGACATCCTCCAGATAGTGTAGCTGAGGAGGCACCT ACACTCACCTGCACTAACAGAGTGGCCGTCCTAACCTCGGGCCTGCTGCGCAGACGTCCATCACGTTAGCTGTCC CACATCACAGACTATGCCATTGGGGTAGTTGTGTTTCAACGGAAAGTGCTGTCTTAAACTAAATGTGCAATAGA AGGTGATGTTGCCATCCTACCGTCTTTTCCTGTTTCCTAGCTGTGTGAATACCTGCTCACGTCAAATGCATACAA GTTTCATTCTCCCTTTCACTAAAAACACACAGGTGCAACAGACTTGAATGCTAGTTATACTTATTTGTATATGGT ATTTATTTTTTCTTTTCTTTACAAACCATTTTGTTATTGACTAACAGGCCAAAGAGTCTCCAGTTTACCCTTCAG GTTGGTTTAATCAATCAGAATTAGAATTAGAGCATGGGAGGGTCATCACTATGACCTAAATTATTTACTGCAAAA AGAAAATCTTTATAAATGTACCAGAGAGAGTTGTTTTAATAACTTATCTATAAACTATAACCTCTCCTTCATGAC AGCCTCCACCCCACAACCCAAAAGGTTTAAGAAATAGAATTATAACTGTAAAGATGTTTATTTCAGGCATTGGAT ATTTTTTACTTTAGAAGCCTGCATAATGTTTCTGGATTTACATACTGTAACATTCAGGAATTCTTGGAGAAGATG GAATGCCTATGGAACTAGTGGATCCAAACTGATCCAGTATAAGACTACTGAATCTGCTACCAAAACAGTTAATCA GTGAGTCGAGTGTTCTATTTTTTTTTTTTTTCCTCCCCTATCTGTATTCCCAAAAATTACTTTGGGGCTAATTT AACAAGAACTTTAAATTGTGTTTTAATTGTAAAAATGGCAGGGGGTGGAATTATTACTCTATACATTCAACAGAG ACTGAATAGATATGAAAGCTGATTTTTTTTAATTACCATGCTTCACAATGTTAAGTTATATGGGGAGCAACAGCA AACAGGTGCTAATTTGTTTTGGATATAGTATAAGCAGTGTCTGTGTTTTGAAAGAATAGAACACAGTTTGTAGTG CCACTGTTGTTTTGGGGGGGGCTTTTTTTCTTTTTCCGGAAAATCCTTAAACCTTAAGATACTAAGGACGTTGTT  $\tt TTGGTTGTACTTGGAATTCTTAGTCACAAAATATATTTTGTTTACAAAAATTTCTGTAAAACAGGTTATAACAGT$ GTTTAAAGTCTCAGTTTCTTGCTTGGGGAACTTGTGTCCCTAATGTGTTAGATTGCTAGATTGCTAAGGAGCTGA TACTTGACAGTTTTTTAGACCTGTGTTACTAAAAAAAAGATGAATGTCGGAAAAGGGTGTTGGGAGGGTGGTCAA CAAAGAAACAAGATGTTATGGTGTTTAGACTTATGGTTGTTAAAAATGTCATCTCAAGTCAAGTCACTGGTCTG TTTGCATTTGATACATTTTTGTACTAACTAGCATTGTAAAATTATTTCATGATTAGAAATTACCTGTGGATATTT GTATAAAAGTGTGAAATAAATTTTTTATAAAAGTGTTCATTGTTTCGTAACACAGCATTGTATATGTGAAGCAAA 

### 369/2825 **FIGURE 337**

MAAAGQLCLLYLSAGLLSRLGAAFNLDTREDNVIRKYGDPGSLFGFSLAMHWQLQPEDKRLLLVGAPRGEALPLQ
RANRTGGLYSCDITARGPCTRIEFDNDADPTSESKEDQWMGVTVQSQGPGGKVVTCAHRYEKRQHVNTKQESRDI
FGRCYVLSQNLRIEDDMDGGDWSFCDGRLRGHEKFGSCQQGVAATFTKDFHYIVFGAPGTYNWKGIVRVEQKNNT
FFDMNIFEDGPYEVGGETEHDESLVPVPANSYLGFSLDSGKGIVSKDEITFVSGAPRANHSGAVVLLKRDMKSAH
LLPEHIFDGEGLASSFGYDVAVVDLNKDGWQDIVIGAPQYFDRDGEVGGAVYVYMNQQGRWNNVKPIRLNGTKDS
MFGIAVKNIGDINQDGYPDIAVGAPYDDLGKVFIYHGSANGINTKPTQVLKGISPYFGYSIAGNMDLDRNSYPDV
AVGSLSDSVTIFRSRPVINIQKTITVTPNRIDLRQKTACGAPSGICLQVKSCFEYTANPAGYNPSISIVGTLEAE
KERRKSGLSSRVQFRNQGSEPKYTQELTLKRQKQKVCMEETLWLQDNIRDKLRPIPITASVEIQEPSSRRRVNSL
PEVLPILNSDEPKTAHIDVHFLKEGCGDDNVCNSNLKLEYKFCTREGNQDKFSYLPIQKGVPELVLKDQKDIALE
ITVTNSPSNPRNPTKDGDDAHEAKLIATFPDTLTYSAYRELRAFPEKQLSCVANQNGSQADCELGNPFKRNSNVT
FYLVLSTTEVTFDTPYLDINLKLETTSNQDNLAPITAKAKVVIELLLSVSGVAKPSQVYFGGTVVGEQAMKSEDE
VGSLIEYEFRVINLGKPLTNLGTATLNIQWPKEISNGKWLLYLVKVESKGLEKVTCEPQKEINSLNLTESHNSRK
KREITEKQIDDNRKFSLFAERKYQTLNCSVNVNCVNIRCPLRGLDSKASLILRSRLWNSTFLEEYSKLNYLDILM
RAFIDVTAAAENIRLPNAGTQVRVTVFPSKTVAQYSGVPWWIILVAILAGILMLALLVFILWKCGFFKRNKKDHY
DATYHKAEIHAQPSDKERLTSDA

# 370/2825 FIGURE 338

# 371/2825 FIGURE 339

 ${\tt MGGFFSSIFSSLFGTREMRILILGLDGAGKTTILYRLQVGEVVTTIPTIGFNVETVTYKNLKFQVWDLGGQTSIR} \\ {\tt PYWRCYYSNTDAVIYVVDSCDRDRIGISKSELVAMLEEEELRKAILVVFANKQDMEQAMTSSEMANSLGLPALKD} \\ {\tt RKWQIFKTSATKGTGLDEAMEWLVETLKSRQ} \\$ 

### 372/2825 FIGURE 340A

GAATTCGTTGTTGGGAAGGACTGGGGAAACAGCTGTAACATTTGCCACCCTCAGAAGCTGCTGGTCCTGTGTCAC ACCACCTTAGCCTCTTGATCGAGGAAGATTCTCGCTGAAGTCTGTTAATTCTACTTTTTTGAGTACTT**ATG**AATAA CCACGTGTCTTCAAAACCATCTACCATGAAGCTAAAACATACCATCAACCCTATTCTTTTATATTTTTATACATTT TCTAATATCACTTTATACTATTTTAACATACATTCCGTTTTATTTTTTCTCCGAGTCAAGACAAGAAAAATCAAA CCGAATTAAAGCAAAGCCTGTAAATTCAAAACCTGATTCTGCATACAGATCTGTTAATAGTTTGGATGGTTTGGC TTCAGTATTATACCCTGGATGTGATACTTTAGATAAAGTTTTTACATATGCAAAAAACAAATTTAAGAACAAAAG ACTCTTGGGAACACGTGAAGTTTTAAATGAGGAAGATGAAGTACAACCAAATGGAAAAATTTTTTAAAAAGGTTAT TCTTGGACAGTATAATTGGCTTTCCTATGAAGATGTCTTTGTTCGAGCCTTTAATTTTGGAAATGGATTACAGAT GTTGGGTCAGAAACCAAAGACCAACATCGCCATCTTCTGTGAGACCAGGGCCGAGTGGATGATAGCTGCACAGGC GTGTTTTATGTATAATTTTCAGCTTGTTACATTATATGCCACTCTAGGAGGTCCAGCCATTGTTCATGCATTAAA CCCACGCCTGCGGCACATCATCACTGTTGATGGAAAGCCACCGACCTGGTCCGACTTCCCCAAGGGCATCATTGT CTCAGATATTGCAGTAATCATGTACACAAGTGGATCCACAGGACTTCCAAAGGGAGTCATGATCTCACATAGTAA CATTATTGCTGGTATAACTGGGATGCCAGAAAGGATTCCAGAACTAGGAGAGGAAGATGTCTACATTGGATATTT ACCACAGACTTTAGCAGATCAGTCTTCAAAAATTAAAAAAGGAAGCAAAGGGGATACATCCATGTTGAAACCAAC ACTGATGGCAGCAGTTCCGGAAATCATGGATCGGATCTACAAAAATGTCATGAATAAAGTCAGTGAAATGAGTAG TTTTCAACGTAATCTGTTTATTCTGGCCTATAATTACAAAATGGAACAGATTTCAAAAGGACGTAATACTCCACT GTGCGACAGCTTTGTTTTCCGGAAAGTTCGAAGCTTGCTAGGGGGGAAATATTCGTCTCCTGTTGTGTGGGGGCGC TCCACTTCTGCAACCACGCAGCGATTCATGAACATCTGTTTCTGCTGTCCTGTTGGTCAGGGATACGGGCTCAC TGAATCTGCTGGGGCTGGAACAATTTCCGAAGTGTGGGACTACAATACTGGCAGAGTGGGAGCACCATTAGTTTG  $\verb|CTGTGAAATCAAATTAAAAAACTGGGAGGAGGTGGATACTTTAATACTGATAAGCCACACCCCAGGGGTGAAAT| \\$ TCTTATTGGGGGCCAAAGTGTGACAATGGGGTACTACAAAAATGAAGCAAAAACAAAAGCTGATTTCTCTGAAGA TCGTAAAAAGGACCTTGTAAAACTACAGGCAGGGGAATATGTTTCTCTTGGGAAAGTAGAGGCAGCTTTGAAGAA TCTTCCACTAGTAGATAACATTTGTGCATATGCAAACAGTTATCATTCTTATGTCATTGGATTTGTTGTGCCAAA TCAAAAGGAACTAACTGAACTAGCTCGAAAGAAAGGACTTAAAGGGACTTGGGAGGAGCTGTGTAACAGTTGTGA AATGGAAAATGAGGTACTTAAAGTGCTTTCCGAAGCTGCTATTTCAGCAAGTCTGGAAAAGTTTGAAATTCCAGT AAAAATTCGTTTGAGTCCTGAACCGTGGACCCCTGAAACTGGTCTGGTGACAGATGCCTTCAAGCTGAAACGCAA  ${\tt AGAGCTTAAAACACATTACCAGGCGGACATTGAGCGAATGTATGGAAGAAAA} {\tt TAAT} {\tt TATTCTCTTCTGGCATCAG}$ TTTGCTACAGTGAGCTCACATCAAATAGGAAAATACTTGAAATGCATGTCTCAAGCTGCAAGGCAAACTCCATTC CTCATATTAAACTATTACTTCTCATGACGTCACCATTTTTAACTGACAGGATTAGTAAAACATTAAGACAGCAAA GAATTTTTCTGAATCATATTGGGGAAGCAGTGATTTTAAAACCTCAAGTTTTTAAACATGATTTATATGTTCTGT ATAATGTTCAGTTTGTAACTTTTTAAAAGTTTGGATGTATAGAGGGATAAATAGGAAATATAAGAATTGGTTATT TGGGGGCTTTTTTACTTACTGTATTTAAAAATACAAGGGTATTGATATGAAATTATGTAAATTTCAAATGCTTAT TTTATTAGAAAAATGAAGTTTGGTTGGTGATGCATGAAACAAAATAGCAAGAGAGGGTTATAGTTTAATAGTAA GGGAGATAACACAGCATGTGTAGCACCAGTTGATAATTGGTCTCTAGTAGCTTACTGTCAAAATGTTCAATGAAG TCTTCTGTTCATCTGTTGAAACTAGGAAAATACCCAAACTTAAATGGAAGAATTCTGAAAGAGAGGATAGAATTT AAAGAACAAGAGTATATAAAGTTATTCTTTGAATATTTCGTTGACTATATGTACATTGAGTTATCTATATTTGTA AACAAATTAGTCATGGAAAATTATTCTATTCCAAAGTCTCCTTTTAGTCTAGATAATCATTATTTCATTTTAAAA TTATCCTTTCTCCCATAACCTTTGTACACTAAAAAAATGAAAGAATTTAGAATGTATTTGATGATAGCATTCTCAC TAAGACACATGAGAATTTAACTTTATAACCGCGTGAGTTAAGATTTAATTCATAGGTTTTGATGTCATTGTTGAA 

# 373/2825 **FIGURE 340B**

## 374/2825 FIGURE 341

MNNHVSSKPSTMKLKHTINPILLYFIHFLISLYTILTYIPFYFFSESRQEKSNRIKAKPVNSKPDSAYRSVNSLD GLASVLYPGCDTLDKVFTYAKNKFKNKRLLGTREVLNEEDEVQPNGKIFKKVILGQYNWLSYEDVFVRAFNFGNG LQMLGQKPKTNIAIFCETRAEWMIAAQACFMYNFQLVTLYATLGGPAIVHALNETEVTNIITSKELLQTKLKDIV SLVPRLRHIITVDGKPPTWSDFPKGIIVHTMAAVEALGAKASMENQPHSKPLPSDIAVIMYTSGSTGLPKGVMIS HSNIIAGITGMAERIPELGEEDVYIGYLPLAHVLELSAELVCLSHGCRIGYSSPQTLADQSSKIKKGSKGDTSML KPTLMAAVPEIMDRIYKNVMNKVSEMSSFQRNLFILAYNYKMEQISKGRNTPLCDSFVFRKVRSLLGGNIRLLLC GGAPLSATTQRFMNICFCCPVGQGYGLTESAGAGTISEVWDYNTGRVGAPLVCCEIKLKNWEEGGYFNTDKPHPR GEILIGGQSVTMGYYKNEAKTKADFSEDENGQRWLCTGDIGEFEPDGCLKIIDRKKDLVKLQAGEYVSLGKVEAA LKNLPLVDNICAYANSYHSYVIGFVVPNQKELTELARKKGLKGTWEELCNSCEMENEVLKVLSEAAISASLEKFE IPVKIRLSPEPWTPETGLVTDAFKLKRKELKTHYQADIERMYGRK

# 375/2825 FIGURE 342

# 376/2825 FIGURE 343

MAPFEPLASGILLLLWLIAPSRACTCVPPHPQTAFCNSDLVIRAKFVGTPEVNQTTLYQRYEIKMTKMYKGFQAL GDAADIRFVYTPAMESVCGYFHRSHNRSEEFLIAGKLQDGLLHITTCSFVAPWNSLSLAQRRGFTKTYTVGCEEC TVFPCLSIPCKLQSGTHCLWTDQLLQGSEKGFQSRHLACLPREPGLCTWQSLRSQIA

### 377/2825 FIGURE 344

 ${\tt CAACCAGGGAGATTTCTCCATTTTCCTCTTGTCTACAGTGCGGCTACAAATCTGGGATTTTTTTATTACTTCTTT}$ TTTTTCGAACTACACTTGGGCTCCTTTTTTTGTGCTCGACTTTTTCCACCCTTTTTCCCTCCTCCTGTGCTGCT GCTTTTTGATCTCTTCGACTAAAATTTTTTTATCCGGAGTGTATTTAATCGGTTCTGTTCTGTCCTCTCCACCAC GTTACACCAACCCGAGGCTCTTTGTTTCCCCTCTTGGATCTGTTGAGTTTCTTTGTTGAAGAAGCCAGC**ATG**GGT GCCCAGTTCTCCAAGACCGCAGCGAAGGGAGAGCCGCCGCGGAGAGGCCTGGGGAGGCGGCTGTGGCCTCGTCG CCTTCCAAAGCGAACGGACAGGAAATGGCCACGTGAAGGTAAACGGCGACGCTTCGCCCGCGGCCGCCGAGTCG GGGGCGCCTCCTCCGCCGAGAAAGGTGAGCCGCCGCCGCCGCTGCCCCCGAGGCCGGGGCCAGCCCG TCGGCCGCCTCCTCGACTTCTTCGCCCAAGGCCGAGGACGGGGCCACGCCCTCGCCCAGCAACGAGACCCCGAAA AAAAAAAAGAGCGCTTTTCCTTCAAGAAGTCTTTCAAGCTGAGCGGCTTCTCCTTCAAGAAGAACAAGAAGGAG GCTGGAGAAGGCGGTGAGGCTGAGGCCCCCCTGCCGAAGGCGGCAAGGACGAGGCCGCCGGGGGCGCAGCTGCG GCCGCCGAGGCGGCGCCTCCGGGGAGCAGCAGCGGCGCGAGGAGGAGCGGCGAGGAGAG GACGAGACCAAGGCCGCGAGGAGCCCAGCAAGGTGGAGGAGAAAAAGGCCGAGGAGGCCGGGGCCAGCGCCCC GCCGCCGCAGCCTCGTCAGCCTGCGCAGCCCCTCACAGGAGGCCCAGCCCGAGTGCAGTCCAGAAGCCCCCCCA TTTTTTTAAGCACCAAATTTTGTTGTTTTTTTTTTTCTCCCCTCCCCACAGATCCCATCTCAAATCATTCTGTT GACATGCCCATATATGAAGGAGATGGGTGGGTCAAAAAGGGATATCAAATGAAGTGATAGGGGTCACAATGGGGA TTAAAGAAAAGTTATTACCATGTATTTTGTGAGGCAGGTTTACAACACTACAAGTCTTGAGTTAAGAAGGAAAGA ATTGTTTATTGCTGGTTTATGACCTTAATAAAGTGTAATTATGTATTACCAGCAGGGTGTTTTTAACTGTGACTA TTGTATAAAAACAAATCTTGATATCCAGAAGCACATGAAGTTTGCAACTTTCCACCCTGCCCATTTTTGTAAAAC TGCAGTCATCTTGGACCTTTTAAAACACAAATTTTAAACTCAACCAAGCTGTGATAAGTGGAATGGTTACTGTTT TCTTCAGATGCAATGGTTTTGTGTAGCATCTTGTCTATCATGTTTTGTAAATACTGGAGAAGCTTTGACCAATTT GACTTAGAGATGGAATGTAACTTTGCTTACAAAAATTGCTATTAAACTCCTGCTTAAGGTGTTCTAATTTTCTGT GAGCACACTAAAAGCGAAAAATAAATGTGAATAAAATGT

# 378/2825 FIGURE 345

MGAQFSKTAAKGEAAAERPGEAAVASSPSKANGQENGHVKVNGDASPAAAESGAKEELQANGSAPAADKEEPAAA GSGAASPSSAEKGEPAAAAAPEAGASPVEKEAPAEGEAAEPGSATAAEGEAASASSTSSPKAEDGATPSPSNET PKKKKKRFSFKKSFKLSGFSFKKNKKEAGEGGEAEAPAAEGGKDEAAGGAAAAAAEAGAASGEQAAAPGEEAAAG EEGAAGGDPQEAKPQEAAVAPEKPPASDETKAAEEPSKVEEKKAEEAGASAAACEAPSAAGPGAPPEQEAAPAEE PAAAAASSACAAPSQEAQPECSPEAPPAEAAE

### 379/2825 FIGURE 346

GTTTGAATGAGACTCGTCGCACCGAAGCCGCCGCCACCACCGCGCCTCCGCCTCGGCCGCCGCAGCTGCTCC CGCTCTACTCCGTTACTGTAAAATGGGGAAAGGAGAAATTTGAAGGTGTAGAATTGAATACAGATGAACCTCCAA CGCTAAAGGATGATGATTGGGGAAACATCAAAATAAAAAATGGAATGACTCTACTAATGATGGGGTCAGCAGATG CTCTTCCAGAAGAACCCTCAGCCAAAACTGTCTTCGTAGAAGAACATGACAGAAGAACAGTTAGCATCTGCTATGG AGTTACCATGTGGATTGACAAACCTTGGTAACACTTGTTACATGAATGCCACAGTTCAGTGTATTCGTTCTGTGC CTGAACTCAAAGATGCCCTTAAAAGGTATGCAGGTGCCTTGAGAGCTTCAGGGGAAATGGCTTCAGCGCAGTATA TTACTGCAGCCCTTAGAGATTTGTTTGATTCCATGGATAAAACTTCTTCCAGTATTCCACCTATTATTCTACTGC AGTTTTTGCACATGGCTTTCCCACAGTTTGCCGAGAAAGGTGAACAAGGACAGTATCTTCAACAGGATGCTAATG AATGTTGGATACAAATGATGCGAGTATTGCAACAGAAATTGGAAGCAATAGAGGATGATTCTGTTAAAGAGACAG ACTCCTCATCTGCATCGGCAGCGACACCTTCTAAAAAGAAAAGTTTAATCGATCAGTTCTTCGGTGTTGAGTTTG AAACTACCATGAAATGTACAGAATCTGAAGAAGAAGAAGTCACCAAAGGAAAGGAAAATCAACTTCAGCTTAGCT GTTTTATCAATCAGGAAGTCAAGTATCTTTTTACAGGACTTAAATTGCGACTTCAGGAAGAAATCACCAAACAGT AGATGGTTCGATTTTTTTATAAAGAGAAGGAATCTGTGAATGCCAAAGTTCTTAAGGATGTTAAATTTCCTCTTA TGTTGGATATGTATGAACTGTGTACACCAGAACTTCAAGAGAAAATGGTGTCTTTTCGATCCAAATTCAAGGATC AACCCTTTTCTTTTGCTGATGATATTTGGCTCCAATAATTGTGGATACTATGACTTACAAGCAGTACTAACACACC AGGGAAGGTCTAGTTCTTCAGGTCATTATGTATCATGGGTGAAAAGGAAACAAGATGAATGGATTAAGTTTGATG ATGACAAAGTCAGCATCGTAACACCAGAAGATATCTTACGGCTTTCTGGTGGTGGAGACTGGCATATCGCTTACG  $\tt TTCTACTCTATGGGCCTCGCAGAGTTGAAATAATGGAAGAGGAAAGTGAACAG\underline{TAA}{\tt TCTTCATTTTAGTATTTAT$ GCTTAGATGTGAAAATAAATGTTATTTGTTGATCATTTCTATAATCCAGAGCTTTAGAGGAAGACACATAGGTGG GTTTATGTTTCACCTCATTTGGAACAAAAGAGGACAGAAGCAGACCACTCTGTGCACCAACCTAAAAAATTACAG AGAAGAAAATTATCTTTGGATTGTGCTGCCCTATATAAAGGTGGCAGAAAGACATTTTTAAAAAAGCTTATTAT TTCTTGCATTATTTTAAAAAGTTCAGAGTTGAAATGCCTTTCAACCATTTCCTTCTGTGGTCATTTTTCTTGCTG CCTTTTCACCCAAGATTCAGCAGTCAGATGTTTACTGCACACCTATTACCTATTATTTGCTGTTCTTGCATGGT TGTTGCTCTCATTGTGTGACTCAGTGCTGCTCCATCCCATGGAAACATGGGCACAATCAAGTATTTGTCCAGC CTATTGCAGGCTTTTCCTGACTTTAAAATAAATTGTGATCAATAATAGTACCTTTGATTATACATTTATTGT GTCTCTCTCTGATGTACTGTGGATTGTACATTTAACTTTGGAATGGCTTTGTAATAATCAGTCTTAAGAAAATGT TGACCTCAAGTCTTTTGTCTTCTGAGTGTTTGGAGCCTTGGCTGAAGACATGTTTAATACTGTACAATTTCTGAAGA TTGGCTCAGTTGTGTTTTCCTTTCTTAATTCCCCAAATCTTGGCTGGGCACGGTGGCTCACGCCTGTAAT  $\verb|CCCAGCACTTTGGGAGGCTGAGGCAGGTTGATTCACTGGAGGTTGAGTTCGAGACCAGCCTGGCCACATGGTGAAA| \\$ CCCTGTCTCTACTAAAGATATAAAAATTAGCTGGATGTGGTGGTGCACACCTGTGATCCCAGCTACTTGGGAGGC TGAGGTGGGAGAATTGCTTGAACCTGGAAGGCAGAAGTTGCAGTGAGCTGAGATCACACCACTGCACTCCAGCCT 

# 380/2825 FIGURE 347

MPLYSVTVKWGKEKFEGVELNTDEPPMVFKAQLFALTGVQPARQKVMVKGGTLKDDDWGNIKIKNGMTLLMMGSA
DALPEEPSAKTVFVEDMTEEQLASAMELPCGLTNLGNTCYMNATVQCIRSVPELKDALKRYAGALRASGEMASAQ
YITAALRDLFDSMDKTSSSIPPIILLQFLHMAFPQFAEKGEQGQYLQQDANECWIQMMRVLQQKLEAIEDDSVKE
TDSSSASAATPSKKKSLIDQFFGVEFETTMKCTESEEEEVTKGKENQLQLSCFINQEVKYLFTGLKLRLQEEITK
QSPTLQRNALYIKSSKISRLPAYLTIQMVRFFYKEKESVNAKVLKDVKFPLMLDMYELCTPELQEKMVSFRSKFK
DLEDKKVNQQPNTSDKKSSPQKEVKYEPFSFADDIGSNNCGYYDLQAVLTHQGRSSSSGHYVSWVKRKQDEWIKF
DDDKVSIVTPEDILRLSGGGDWHIAYVLLYGPRRVEIMEEESEQ

### 381/2825 FIGURE 348

GAGGAGCTCTGCGCGCGCGGCGGCGGCGATCCGAGCCGGGACGGGCTGCAGGGGGGGTGCTGCAGAGGACACGAG TGTTGCTGCCACGATCAGTGCCACAGAGACCCTCTCGGAAGAGGAGCAGGAAGAGCTAAGAAGAGAACTTGCAAA GGTAGAAGAAGAATCCAGACTCTGTCTCAAGTGTTAGCAGCAAAAGAGAAGCATCTAGCAGAGATCAAGCGGAA ACTTGGAATCAATTCTCTACAGGAACTAAAACAGAACATTGCCAAAGGGTGGCAAGACGTGACAGCAACATCTGC TTACAAGAAGACATCTGAAACCTTATCCCAGGCTGGACAGAAGGCCTCAGCTGCTTTTTCGTCTGTTGGCTCAGT CATCACCAAAAAGCTGGAAGATGTAAAAAACTCCCCAACTTTTAAATCATTTGAAGAAAAGGTCGAAAACTTAAA GTCTAAAGTAGGGGGAACCAAGCCTGCTGGTGGTGATTTTGGAGAAGTCTTGAATTCGGCTGCAAATGCTAGTGC CACCACCACGGAGCCTCTTCCAGAAAAGACACAGGAGAGCCTG**TGA**GATTCCTACCTTTGTTCTGCTACCCACTG CCAGATGCTGCAAGCGAGGTCCAAGCACATCTTGTCAACATGCATTGCCATGAATTTCTACCAGATGTGCTTTTA TTTAGCTTTACATATTCCTTTGACCAAATAGTTTGTGGGTTAAACAAAATGAAAATATCTTCACCTCTATTCTTG GGAAACACCCTTTAGTGTACATTTATGTTCCTTTATTTAGGAAACACCATTATAAAAACACTTATAGTAAATGGG GACATTCACTATAATGATCTAAGAAGCTACAGATTGTCATAGTTGTTTTCCTGCTTTACAAAATTGCTCCAGATC TGGAATGCCAGTTTGACCTTTGTCTTCTATAATATTTCCTTTTTTTCCCCTCTTTGAATCTCTGTATATTTGATT CTTAACTAAAATTGTTCTCTTAAATATTCTGAATCCTGGTAATTAAAAGTTTGGGTGTATTTTCTTTACCTCCAA GGAAAGAACTACTAGCTACAAAAAATATTTTGGAATAAGCATTGTTTTGGTATAAGGTACATATTTTGGTTGAAG AGGTGACTTGAGAGTGGAACCTCCTATATCATTATTTAGCACCGTTTGTGACAGTAACCATTTCAGTGTATTGTT TATTATACCACTTATATCAACTTATTTTTCACCAGGTTAAAATTTTAATTTCTACAAAATAACATTCTGAATCAA GCACACTGTATGTTCAGTAGGTTGAACTATGAACACTGTCATCAATGTTCAGTTCAAAAGCCTGAAAGTTTAGAT CTAGAAGCTGGTAAAAATGACAATATCAATCACATTAGGGGAACCATTGTTGTCTTCACTTAATCCATTTAGCAC TATTTAAAATAAGCACACCAAGTTATATGACTAATATAACTTGAAAATTTTTTATACTGAGGGGTTGGTGATAAC TCTTGAGGATGTAATGCATTAATAAAAATCAACTCATCATTTTCTACTTGTTTTCAATGTGTTGGAAACTGTAAA ATGATACTGTAGAACCTGTCTCCTACTTTGAAAACTGAATGTCAGGGCTGAGTGAATCAAAGTGTCTAGACATAT TTGCATAGAGGCCAAGGTATTCTATTCTAATAACTGCTTACTCAACACTACCACCTTTTCCTTATACTGTATATG TGCCAAGCTTGGAGTGACCTAAAGCATTTTTTAAAAGCATGGCTAGATTTACTTCAGTATAAATTATCTTATGAA AACCAAATTTTAAAAGCCACAGGTGTTGATTGTTATAAAATAACATGCTGCCATTCTTGATTGCTAGAGTTTTTG TTAGTACTTTGGATGCAATTAAAACTATGTGCTATCACATGTGAAAAGCTTAATAAATTCCATCTATCAGTAGTA TAGGTCTCAATATTTATTATGAGACCAGTGGTCTGGAAACAGCTTGTTGTACCGAATCAACTGGAGTCTATGCTT AAAAAAAAATTTTTTTTTAACCATCCTTAAATTATTGCTTAATGGTATCATATTAACATATTCTAAATAAGGG GTGCTTACTCAAGTGTTAAATTATTTTTTCTATGAACTAGTCTACTTCTTAAAATTCAAACATATTCTTTTGATC ACATTGTTTCTTGAGCATCCTGCCCTGCTACTAACTTTTCAACAAGGCAAAATGGAGTAAAGTGGCAATTTCTTT AGATGAGTGAAATACCCTCAAGTCTCTTTTCTGCCCAAAAAGGGGAAAAGTGATAGAAATGGGGGTGGCAAGTGGG TATACTTAATCTAAGAAACAAAGTAATCTACTGTAAAGTACTCTGCCCCTTGAAAGAAGTATTAAAAAGAGTGAG GATGGATTTAGAAAAAACATGAATTTAGAAATATTCAAAATGGTTTTTGTGGCAGATTCAATATTATGAATTCA CAGATATTTAAAGAATGAGAAACATAGTAATTAGTAGAAATGCCAGAAACAGTTCCTGGTTCCTCTTGTGTTTGA CACTAAGAAATAGCAAGAGTGTGAAATCTCAGATACTTATGAAATCTCACAGATGTAAGGACTCAAGTGTAGAA GAAAATATCCCCTTCTTACAAAAAGAAATGTCAATTTATGGAGTTTGTGGGAAATAGGGCAAGAATTCTTATGCT TATGAGAGCCAAGTAGTCAGTGGAAGAGAGTAGAGCTCAAAACTGGATTATCACCTTAGCAACTTAGAATAGTTT GAAATAGAAAAAAGTATTTAATTTGGATCTGGATCTGTTAAGATATGCACAGTCTATTTTTTTGTATAGTATTGG AAAATAAAAATGCTATAATTTG

## 382/2825 FIGURE 349

MDRGEQGLLRTDPVPEEGEDVAATISATETLSEEEQEELRRELAKVEEEIQTLSQVLAAKEKHLAEIKRKLGINS LQELKQNIAKGWQDVTATSAYKKTSETLSQAGQKASAAFSSVGSVITKKLEDVKNSPTFKSFEEKVENLKSKVGG TKPAGGDFGEVLNSAANASATTTEPLPEKTQESL

### 383/2825 FIGURE 350A

AGGTAGCAAGATGGCCGCCGCTGAGGAAGGCTGTAGTGTCGGGGCCGAAGCGGACAGGGAATTGGAGGAGCTTCT GGAAAGTGCTCTTGATGATTTCGATAAAGCCAAACCCTCCCCAGCACCCCTTCTACCACCACGGCCCCTGATGC GGAACTATTCGACAGTGAACTGGCTTCCCAAGCCACTGCGGAGTTCGAGAAGGCAATGAAGGAGTTGGCTGAGGA AGAACCCCACCTGGTGGAGCAGTTCCAAAAGCTCTCAGAGGCTGCAGGGAGAGTGGGCAGTGATATGACCTCCCA ACAAGAATTCACTTCTTGCCTAAAGGAAACACTAAGTGGATTAGCCAAAAAATGCCACTGACCTTCAGAACTCCAG CATGTCGGAAGAGCTGACCAAGGCCATGGAGGGGCTAGGCATGGACGAAGGGGATGGGGAAGGGAACATCCT CCCCATCATGCAGAGTATTATGCAGAACCTACTCTCCAAGGATGTGCTGTACCCATCACTGAAGGAGATCACAGA AAAGTATCCAGAATGGTTGCAGAGTCATCGGGAATCTCTACCTCCAGAGCAGTTTGAAAAATATCAGGAGCAGCA GCCTCCTGGCCTCAACTTTGACCTGGATGCCCTCAATCTTTCGGGCCCACCAGGTGCCAGTGGTGAACAGTGTCT GATCATG**TGA**AACACACGTTTTCCTCTGAGTCCCAGCTATGGGGAACATCTGGAGTCAGCAGAACCATTG GGACCTGAGGCAGGAGTGTCACCTGCGGGAGAAGTCTGCCCGCTGCCCTCTGTCATCCCATTCAAGATTGTGCCA TACCAGCTGAGGTTTTTCCTCTGTCTCTCTAGGAATAGGGTCTGTTTCACAGGCCATTTCTGTGAACCCTACTCC TTCTTTCCTTTCTCAAAATAATGTTATATGTGGCCACACTGATGTTCACCTTTACGTCCAGGGTCTTTGTGCCTT GTCTCTACTCCTCTTTGGATCTGGGGAGGGGGGGAGAGACCTGGGACTCTGTATTTCTATAGTTCTCCTGGC AGAGCCTTTGAGAATGGGGAGAAACAGCCTGGGCTGGGGCTACAGGTCTGTCACTATGCTCTCTTGCCTTCAGAC AGACCATTCTGAATTCTCTAAAGGGAAAGGGCTTTTGCATCTAATCACAATAGAGTTGAAAGAGAGGCCTTAGGA TTCTCCTCTCTAGGTGCTGAGCCCTCACCTCCCTGTTCCAGGCTGAGAACTCAAATGGTTACCCTGCTTCTTC  $\tt CTACAATGCTGTGATATGGGTGAACCCAGCCCCTGACCTTCCTCTATCCCCTGCCCATCCTCCCTTTTACCTC$ CTCTCTTTTTAAACACCTGTTTATCCCAACCTTTTTGAGCTCAAGCTGTGATAAAGAAGGGGCCCATCCTATTTC CCCTCATCTAGTCCATTTACGATTCTCACTGACTCCCCGTCTTCCTGGCAGACACAAATAAACCCAGTGTCAGGT CTAGGAAATTAATGGCTATTCTTCCCCAGATACATTCTGGCTTATTTGAGATACATGATTCTCTTAGAATCCTGT CCCTTGGTTCAGGAAAGTAGCTTGGAAAAGGAGTAGGGGTATAGCTTGGGTCCCTTTTCCTGCAAGGCCCCATGG GGCAGAATATAATAATATTCTGAGTGAGGAGTGTGGTCTTTTTCTGATCTTCCTCAGCTTCCGTAAGTTGCAGA AGTCTGCACCTCAGTTGGCAAAGGAGACTGGATGGCCATCTTTCCTCATGTTCCCTTGAGTATTTCAATGTAGAA AGCCCTTCAAGTGGTATTATATTTTAACCTTTTACATTATTGTTATTAATGTTAGTAATATATTGTTATGTTTTC TAAATTATTTTTTTTAAGCTGACGTGGCTTTTTTTCTGTGGCTCCCAGTGGGTCTACGGACCTTGGCTGACATA TGTTGGTAGGTACTCTGGTCAGCTGGCTGTCCTGGTTCACTCAGAAGATAAGTCTCTCCAAAGCAAATTC ACATGCATTATGAGTCGCTTTGAGCTTCTGACATGTCACTTGCCCCGAGGTTAAAACTTTTCACCCCTTGAAGAC CTTACATGTTTTATGGTATTGGTGAGGAAGGAAATGTTCTCAAGGTCTCAGGCTATTTGGGAAATTCCAACTCCT ATACCTTACCAGAGCATGGAAGAGCCCAGATCTGAATGTAAAACGTCTCTGTTCTGCCAGAGATGGAAAAAATAC AGGTATACTTGTGATATAGTCATGGGGCTTCAGTGTCACTATTTTCTCCTTAAAGCTCCAGCCAAAAACTGGACA AGGATAGAGAGGAGGAGGAAGAACAAAAGAGCCCTTCTCTATGAACCTTGTGCCTTCTGTCCTACCAGTTTTCT TTTACAGATTCTCACTTCTGCTAGCCTAGCCAGGGCTTACTCCAGGAATCTAAATAGATGCCCTAGTCCACTTTA TTCCAAAAACTTTTGGAATTATTCCAGGACTTGTGGTGGAGTTATGGTACTCTAGGGCAGTCTTTCTCAAACTAT GTATGGTAAAGGACCAGGTTTTTTGTTTTCCAGTCCTTCACTTATCAATATGCATTCCTATTGCCGATGACAGGT ATGGAGTTCACACTGTGTGCCGCCGGCCAAGTTTGACAGCACCCAAACTGGCCAGACTGTTCTGTAGGTTA AGTCCATTGATCATGTACTTGGATATCACAGCAACATTGAAATGCTAAAAAGTTTTTAAACACTCTCAATTTCTA ATTCACCATGTCACAGACTGGTGAAAAAAAAAAAAAGGTGTTCACTGACCAGCACAAGTCTGCAGATCATCTTTG CTCCTATTTGAGTTTCACTTCCAGAGAACTTGTTCTTCAGCAAGAATGTGTCACTAGTAAGGACATCTCTAGCAT TTCTCTAGCCTTCCTTTTCTGCTGCTCAAAAATAATCGTTACAAAGCTTAGGTTTAAGCTGTATATGAAATATTT ATGCGACTCTCAAACTTTAAAGGAGTTGCTCCTTTGTTCCAAAATTAAATGTGTTAGATAAATTTGTGATTGTAT

## 384/2825 FIGURE 350B

## 385/2825 FIGURE 351

MAAAEEGCSVGAEADRELEELLESALDDFDKAKPSPAPPSTTTAPDASGPQKRSPGDTAKDALFASQEKFFQELF DSELASQATAEFEKAMKELAEEEPHLVEQFQKLSEAAGRVGSDMTSQQEFTSCLKETLSGLAKNATDLQNSSMSE EELTKAMEGLGMDEGDGEGNILPIMQSIMQNLLSKDVLYPSLKEITEKYPEWLQSHRESLPPEQFEKYQEQHSVM CKICEQFEAETPTDSETTQKARFEMVLDLMQQLQDLGHPPKELAGEMPPGLNFDLDALNLSGPPGASGEQCLIM

### 386/2825 FIGURE 352

GGGTTCGAGTTCTGGAAACAGCTGTGCGCCGAGCATGGTATCAGCCCCGAGGCGATCGTGGAGGAGTTCGCCACC GAGGGCACTGACCGCAAGGACGTCTTTTTCTACCAGGCAGACGATGAGCACTACATCCCCCGGGCCGTGCTGCTG GACTTGGAACCCCGGGTGATCCACTCCATCCTCAACTCCCCCTATGCCAAGCTCTACAACCCAGAGAACATCTAC CTGTCGGAACATGGAGGAGCTGGCAACAACTGGGCCAGCGGATTCTCCCAGGGAGAAAAGATCCATGAGGAC ATTTTTGACATCATAGACCGGGAGGCAGATGGTAGTGACAGTCTAGAGGGCTTTGTGCTGTGTCACTCCATTGCT GGGGGGACAGGCTCTGGACTGGGTTCCTACCTCTTAGAACGGCTGAATGACAGGTATCCTAAGAAGCTGGTGCAG ACATACTCAGTGTTTCCCAACCAGGACGAGATGAGCGATGTGGTGGTCCAGCCTTACAATTCACTCCTCACACTC AAGAGGCTGACGCAGAATGCAGACTGTCTGGTGGTGCTGGACAACACACCCCTGAACCGGATTGCCACAGACCGC GTCATGAGGCGGCTGCTGCAGCCCAAGAACGTGATGGTGTCCACAGGCCGAGACCGCCAGACCACCGCTAC  $\verb|CCCTCGGCCCACCGGGTCAGCGGGCTCATGATGGCCAACCACCAGCATCTCCTCGCTCTTCGAGAGAACCTGT| \\$ CGCCAGTATGACAAGCTGCGTAAGCGGGAGGCCTTCCTGGAGCAGTTCCGCAAGGAGACATGTTCAAGGACAAC AGCCCTGCCTGACTGACCACCCCCTCAGAGCACAGATCAGGGACCTCACGCATCTCTTTCTCATATACATGGACT CTCTGTTGGCCTGCAAACACATTTACTTCTCCTCTTATGAGACTATTTATCTTTAATAAAGCACTGGG

## 387/2825 FIGURE 353

MPREIITLQLGQCGNQIGFEFWKQLCAEHGISPEAIVEEFATEGTDRKDVFFYQADDEHYIPRAVLLDLEPRVIH SILNSPYAKLYNPENIYLSEHGGGAGNNWASGFSQGEKIHEDIFDIIDREADGSDSLEGFVLCHSIAGGTGSGLG SYLLERLNDRYPKKLVQTYSVFPNQDEMSDVVVQPYNSLLTLKRLTQNADCLVVLDNTALNRIATDRLHIQNPSF SQINQLVSTIMSASTTTLRYPGYMNNDLIGLIASLIPTPRLHFLMTGYTPLTTDQSVASVRKTTVLDVMRRLLQP KNVMVSTGRDRQTNHCYIAILNIIQGEVDPTQVHKSLQRIRERKLANFIPWGPASIQVALSRKSPYLPSAHRVSG LMMANHTSISSLFERTCRQYDKLRKREAFLEQFRKEDMFKDNFDEMDTSREIVQQLIDEYHAATRPDYISWGTQE O

#### 388/2825 FIGURE 354A

 $\tt TTTTTAAATTTTGCATTTGACTTAAAGTGCC \textbf{ATG} AGAAAATTTGCATACTGCAAGGTGGTCCTAGCCACCTCCTT$ AGGACTTCCTGCTGGAGATGTTCTAGAGCCAGTACAAAAGCCTCATGAAGGTCCTGGAGAAATGGGGAAACCAGT GATGATTGCACTCAACAGATCTTTACCAGATGTTAGGTTAGAAGGGTGTAAAACAAAGGTGTATCCAGATAATCT TCCTACAACAAGTGTGGTGATTGTTTTCCACAATGAGGCTTGGAGCACCTTCTGCGAACTGTCCATAGTGTCAT TAATCGCTCACCAAGACACATGATAGAAGAAATTGTTCTAGTAGATGATGCCAGTGAAAGAGACTTTTTGAAAAG GCCTTTAGAGAGTTATGTGAAAAAACTAAAAGTACCAGTTCATGTAATTCGAATGGAACAACGTTCTGGATTGAT CAGAGCTAGATTAAAAGGAGCTGCTGTGTCTAAAGGCCAAGTGATCACCTTCCTGGATGCCCATTGTGAGTGTAC AGTGGGATGGCTGGAGCCTCTCTTGGCCAGGATCAAACATGACAGGAGAACAGTGGTGTCTCCCATCATCGATGT GATCAGTGATGATACTTTTGAGTACATGGCAGGCTCTGATATGACCTATGGTGGGTTCAACTGGAAGCTCAATTT TCGCTGGTATCCTGTTCCCCAAAGAGAAATGGACAGAAGGAAAGGTGATCGGACTCTTCCTGTCAGGACACCTAC CATGGCAGGAGGCCTTTTTTCAATAGACAGAGATTACTTTCAGGAAATTGGAACATATGATGCTGGAATGGATAT TTGGGGAGGAGAAACCTAGAAATTTCCTTTAGGATTTGGCAGTGTGGAGGAACTTTTGGAAATTGTTACATGCTC ACATGTTGGACATGTGTTTCGGAAAGCTACACCTTACACGTTTCCAGGAGGCACAGGGCAGATTATCAATAAAA TAACAGACGACTTGCAGAAGTGTGGATGGATGAATTCAAGAATTTCTTCTATATAATTTCTCCAGGTGTTACAAA GGTAGATTATGGAGATATATCGTCAAGAGTTGGTCTAAGACAAACTACAATGCAAACCTTTTTCCTGGTACCT AGAGAATATATCCTGATTCTCAAATTCCACGTCACTATTTCTCATTGGGAGAGATACGAAATGTGGAAACGAA TCAGTGTCTAGATAACATGGCTAGAAAAGAGAATGAAAAAGTTGGAATTTTTAATTGCCATGGTATGGGGGGTAA TCAGGTTTTCTCTTATACTGCCAACAAAGAAATTAGAACAGATGACCTTTGCTTGGATGTTTCCAAACTTAATGG CCCAGTTACAATGCTCAAATGCCACCACCTAAAAGGCAACCCAACTCTGGGAGTATGACCCAGTGAAATTAACCCT GCAGCATGTGAACAGTAATCAGTGCCTGGATAAAGCCACAGAAGAGGATAGCCAGGTGCCCAGCATTAGAGACTG CAATGGAAGTCGGTCCCAGCAGTGGCTTCTTCGAAACGTCACCCTGCCAGAAATATTC**TGA**GACCAAATTTACAA AAAAACGAAAAAATAAGGATTGACTGGGCTACCTCAGCATACATTTCTGCCACATTCTTAAGTAGCAAAAAAGG AAAAGTGCTTTCCTCCTCTGCAGGATGTAAGGTTTATCAGCCATTAAAACTTAGACTTCTCTAGCTTTTCACTAG CTGTGAACCAGCCTTCCTGTCCATGGACGTGAAACTGCATAGTAATGAGACTGTGCACACTGATGTTTACAAGAT GGAGAACTGTACCTTCTATGGTTTGCTTGCACATCAGTAGTTTCTGCTGAACGTGCTGTCATAATGAAGAGATTT CCAAGATTTTTTTCCTGATTAGAACGGGTAGCCAGTATATTAAATATTGATAGAAAAATAAAAGAACTGGAACC AAAACAGAACTATGAAGAAGTACAATTTGTTATAGTATAGTATCAAATTTCTATATAGATTTTATACCTCAGTGG GGAAAAATAACTGATTCCAATGACATTCATTTTGTTTTCATCTGTGATAGTCATGGATGCTTTTATTTTCCTTGG GTTTGTGGGCTGTTGGAATTGTAATTTTTAATTGCCTTCTAAAAAATGGAAATTTAACAATGTCTGATCTCAGCT TTATCACATTTTCGACTTCTTCACTTGACCTAACTGATTATGCGAAATACCCAAGATTCATGCTACTGTTCCACA TTTGTTTTCACAGCAATAAATCTTCAGTTCTGTTGTTTATGATTCCACTTAACAAGGGGCCTGCAAATGTGATTT ATTATTTGGGTATTTGGAGATAATACATTTGAGGGTTTTTTTGGAAAACCTTTTTCACTCCATACTCAAATATGCT ACTGAAAATGTTTAATTGGCATTTTTTAATGACTTACCCAAGAAAAGTGCAGCTATTATTCCATATTAATAGGCT TCACCTGAGTGTCAATTGAAAAGTTGTCAATTAAAAGGTAACCTTTTAACTCTCGTAGGAGGAATCTCATTAAGA AAAAGTCGCATCGCTAATGATATTTGCCAAGTTGAGTGTACACAAAGTTTCTCATATCCTGTTCAAGTTAATCAA CATCAAACACATGGGGATGCTTTAGGGTGAGTCTATAATACAAAATGCATAAACCATGTCCCCAGGAAATTTGAA AGGAAGCAAGTGCTGAATGGAATTTTTTTCCTTTTCCATGAGCTGTGTTAATTCTATCTCCAGTAGGCCTAATGC TTGAAATAAGCAAGATGTCTAATCAATAAATTATTTTCATGCTCAGAATTTCAGGTTTTTGTACTCCAGCATAGC

## 389/2825 FIGURE 354B

### 390/2825 FIGURE 355

MRKFAYCKVVLATSLIWVLLDMFLLLYFSECNKCDEKKERGLPAGDVLEPVQKPHEGPGEMGKPVVIPKEDQEKM KEMFKINQFNLMASEMIALNRSLPDVRLEGCKTKVYPDNLPTTSVVIVFHNEAWSTLLRTVHSVINRSPRHMIEE IVLVDDASERDFLKRPLESYVKKLKVPVHVIRMEQRSGLIRARLKGAAVSKGQVITFLDAHCECTVGWLEPLLAR IKHDRRTVVCPIIDVISDDTFEYMAGSDMTYGGFNWKLNFRWYPVPQREMDRRKGDRTLPVRTPTMAGGLFSIDR DYFQEIGTYDAGMDIWGGENLEISFRIWQCGGTLEIVTCSHVGHVFRKATPYTFPGGTGQIINKNNRRLAEVWMD EFKNFFYIISPGVTKVDYGDISSRVGLRHKLQCKPFSWYLENIYPDSQIPRHYFSLGEIRNVETNQCLDNMARKE NEKVGIFNCHGMGGNQVFSYTANKEIRTDDLCLDVSKLNGPVTMLKCHHLKGNQLWEYDPVKLTLQHVNSNQCLD KATEEDSQVPSIRDCNGSRSQQWLLRNVTLPEIF

### 391/2825 FIGURE 356

TGCTGCCGCTGGTGCACGTCTCTGCGACCACGCCAGAACCTTGTGAGCTGGACGATGAAGATTTCCGCTGCGTCT CCGGCGGTCTCAACCTAGAGCCGTTTCTAAAGCGCGTCGATGCGGACGCCGACCCGCGGCAGTATGCTGACACGG  ${\tt TCAAGGCTCTCCGCGTGCGGCGCTCACAGTGGGAGCCGCACAGGTTCCTGCTCAGCTACTGGTAGGCGCCCTGC}$ GTGTGCTAGCGTACTCCCGCCTCAAGGAACTGACGCTCGAGGACCTAAAGATAACCGGCACCATGCCTCCGCTGC CTCTGGAAGCCACAGGACTTGCACTTTCCAGCTTGCGCCTACGCAACGTGTCGTGGGCGACAGGGCGTTCTTGGC TCGCCGAGCTGCAGCAGTGGCTCAAGCCAGGCCTCAAGGTACTGAGCATTGCCCAAGCACACTCGCCTTCTT  $\verb|CCTGCGAACAGGTTCGCGCCTTCCCGGCCCTTACCAGCCTAGACCTGTCTGACAATCCTGGACTGGGCGAACGCG|\\$ GACTGATGGCGGCTCTCTGTCCCCACAAGTTCCCGGCCATCCAGAATCTAGCGCTGCGCAACACAGGAATGGAGA CGCCCACAGGCGTGTGCGCCGCACTGGCGGCGGCAGGTGTGCAGCCCCACAGCCTAGACCTCAGCCACAACTCGC TGCGCGCCACCGTAAACCCTAGCGCTCCGAGATGCATGTGGTCCAGCGCCCTGAACTCCCTCAATCTGTCGTTCG CTGGGCTGGAACAGGTGCCTAAAGGACTGCCAGCCAAGCTCAGAGTGCTCGATCTCAGCTGCAACAGACTGAACA GGGCGCCGCAGCCTGACGAGCTGCCCGAGGTGGATAACCTGACACTGGACGGGAATCCCTTCCTGGTCCCTGGAA CTGCCCTCCCCACGAGGGCTCAATGAACTCCGGCGTGGTCCCAGCCTGTGCACGTTCGACCCTGTCGGTGGGGG  $\tt TGTCGGGAACCCTGGTGCTCCAAGGGGCCCGGGGCTTTGCC\textbf{TAA} GATCCAAGACAGAATAATGAATGGACTC$ AAACTGCCTTGGCTTCAGGGGAGTCCCGTCAGGACGTTGAGGACTTTTCGACCAATTCAACCCTTTGCCCCACCT TTATTAAAATCTTAAAC

## 392/2825 FIGURE 357

MERASCLLLLLLPLVHVSATTPEPCELDDEDFRCVCNFSEPQPDWSEAFQCVSAVEVEIHAGGLNLEPFLKRVDA DADPRQYADTVKALRVRRLTVGAAQVPAQLLVGALRVLAYSRLKELTLEDLKITGTMPPLPLEATGLALSSLRLR NVSWATGRSWLAELQQWLKPGLKVLSIAQAHSPAFSCEQVRAFPALTSLDLSDNPGLGERGLMAALCPHKFPAIQ NLALRNTGMETPTGVCAALAAAGVQPHSLDLSHNSLRATVNPSAPRCMWSSALNSLNLSFAGLEQVPKGLPAKLR VLDLSCNRLNRAPQPDELPEVDNLTLDGNPFLVPGTALPHEGSMNSGVVPACARSTLSVGVSGTLVLLQGARGFA

### 393/2825 FIGURE 358

GGCATCCAAGCAAGTGAAGATGTTAAAGAGATCTTTGCCAGAGCCAGAAATGGAAAGTACAGACTTCTGAAAATA TCTATTGAAAATGAGCAACTTGTGATTGGATCATATAGTCAGCCTTCAGATTCCTGGGATAAGGATTATGATTCC TTTGTTTTACCCCTGTTGGAGGACAAACAACCATGCTATATATTATTCAGGTTAGATTCTCAGAATGCCCAGGGA TATGAATGGATATTCATTGCATGGTCTCCAGATCATTCTCATGTTCGTCAAAAAATGTTGTATGCAGCAACAAGA GCAACTCTGAAGAAGGAATTTGGAGGTGGCCACATTAAAGATGAAGTATTTGGAACAGTAAAGGAAGATGTATCA CAGATTAAAATCAATGAGGTACAGACTGACGTGGGTGTGGACACTAAGCATCAAACACTACAAGGAGTAGCATTT CCCATTTCTCGAGAAGCCTTTCAGGCTTTGGAAAAATTGAATAATAGACAGCTCAACTATGTGCAGTTGGAAATA AAGGATTCAGCTCGTTACCATTTCTTTCTGTATAAACATTCCCATGAAGGAGACTATTTAGAGTCCATAGTTTTT ATTTATTCAATGCCTGGATACACATGCAGTATAAGAGAGCGGATGCTGTATTCTAGCTGCAAGAGCCGTCTGCTA GAAATTGTAGAAAGACAACTACAAATGGATGTAATTAGAAAGATCGAGATAGACAATGGGGATGAGTTGACTGCA GACTTCCTTTATGAAGAAGTACATCCCAAGCAGCATGCACAAGCAAAGTTTTTGCAAAACCAAAAGGTCCTGCA GGAAAAAGAGGAATTCGAAGACTAATTAGGGGCCCAGCGGAAACTGAAGCTACTACTGAT**TAA**AGTCATCACATT AAACATTGTAATACTAGTTTTTTAAAAGTCCAGCTTTTAGTACAGGAGAACTGAAATCATTCCATGTTGATATAA AGTAGGGAAAAAATTGTACTTTTTGGAAAATAGCACTTTTCACTTCTGTGTGTTTTTTAAAATTAATGTTATAGA AGACTCATGATTTCTATTTTTGAGTTAAAGCTAGAAAAGGGTTCAACATAATGTTTAATTTTTGTCACACTGTTTT CATAGCGTTGATTCCACACTTCAAATACTTCTTAAAATTTTATACAGTTGGGCCAGTTCTAGAAAGTCTGATGTC TCAAAGGGTAAACTTACTACTTTCTTGTGGGACAGAAAGACCTTAAAATATTCATATTACTTAATGAATATGTTA ÁGGACCAGGCTAGAGTATTTTCTAAGCTGGAAACTTAGTGTGCCTTGGAAAAGCCGCAAGTTGCTTACTCCGAGT AGCTGTGCTAGCTCTGTCAGACTGTAGGATCATGTCTGCAACTTTTAGAAATAGTGCTTTATATTGCAGCAGTCT TTTATATTTGACTTTTTTTAATAGCATTAAAATTGCAGATCAGCTCACTCTGAAACTTTAAGGGTACCAGATAT TTTCTATACTGCAGGATTTCTGATGACATTGAAAGACTTTAAACAGCCTTAGTAAATTATCTTTCTAATGCTCTG TGAGGCCAAACATTTATGTTCAGATTGAAATTTAAATTAATATCATTCAAAAGGAAACAAAAAATGTTGAGTTTT AAAAATCAGGATTGACTTTTTCTCCAAAACCATACATTTATGGGCAAATTGTGTTCTTTATCACTTCCGAGCAA ATACTCAGATTTAAAATTACTTTAAAGTCCTGGTACTTAACAGGCTAACGTAGATAAACACCTTAATAATCTCAG TTAATACTGTATTTCAAAACACATTTAACTGTTTTCTAATGCTTTGCATTATCAGTTACAACCTAGAGAGATTTT GAGCCTCATATTTCTTTGATACTTGAAATAGAGGGAGCTAGAACACTTAATGTTTAATCTGTTAAACCTGCTGCA AGAGCCATAACTTTGAGGCATTTTCTAAATGAACTGTGGGGGATCCAGGATTTGTAATTTCTTGATCTAAACTTTA TGCTGCATAAATCACTTATCGGAAATGCACATTTCATAGTGTGAAGCACTCATTTCTAAACCTTATTATCTAAGG TAAAATAGAATTTAGAGTATTTGGGGTTTTGTTTGTTTACAAATAATCAGACTATAATATTTAAACATGCAAAAT TGTCATGAAGGTATAAAATAATTTAGAAGTGAATGTTTTTCTGTACCATCTATGTGCAATTATACTCTAAATTCC ACTACACTACATTAAAGTAAATGGACATTCCAGAATATAGATGTGATTATAGTCTTAAACTAATTATTATTAAAC CAATGATTGCTGAAAATCAGTGATGCATTTGTTATAGAGTATAACTCATCGTTTACAGTATGTTTTAGTTGGCAG TATCATACCTAGATGGTGAATAACATATTCCCAGTAAATTTATATAGCAGTGAAGAATTACATGCCTTCTGGTGG 

## 394/2825 FIGURE 359

MSHQTGIQASEDVKEIFARARNGKYRLLKISIENEQLVIGSYSQPSDSWDKDYDSFVLPLLEDKQPCYILFRLDS QNAQGYEWIFIAWSPDHSHVRQKMLYAATRATLKKEFGGGHIKDEVFGTVKEDVSLHGYKKYLLSQSSPAPLTAA EEELRQIKINEVQTDVGVDTKHQTLQGVAFPISREAFQALEKLNNRQLNYVQLEIDIKNEIIILANTTNTELKDL PKRIPKDSARYHFFLYKHSHEGDYLESIVFIYSMPGYTCSIRERMLYSSCKSRLLEIVERQLQMDVIRKIEIDNG DELTADFLYEEVHPKQHAHKQSFAKPKGPAGKRGIRRLIRGPAETEATTD

### 395/2825 FIGURE 360

## 396/2825 FIGURE 361

 ${\tt MPFLDIQKRFGLNIDRWLTIQSGEQPYKMAGRCHAFEKEWIECAHGIGYTRAEKECKIEYDDFVECLLRQKTMRR} \\ {\tt AGTIRKQRDKLIKEGKYTPPPHHIGKGEPRP}$ 

## 397/2825 FIGURE 362

CGGACGCGTGGGGCCCGGGGGGCTGAGGCGCCGCCGCCGCGGGGGGCCGCTCGCGTCCTCCATCGAGGCCG GAGAGGAACCGCTGCTGCCGAACTCAAGCCCGGGCGCCCCCACCAGTTTGATTGGAAGTCCAGCTGTGAAA CCTGGAGCGTCGCCTTCTCCCCAGATGGCTCCTGGTTTGCTTGGTCTCAAGGACACTGCATCGTCAAACTGATCC CCTGGCCGTTGGAGGAGCAGTTCATCCCTAAAGGGTTTGAAGCCAAAAGCCGAAGTAGCAAAAATGAGACGAAAG GGCGGGCCAAAAGAGAGACGCTGGACTGTGGTCAGATTGTCTGGGGGCTGGCCTTCAGCCCGTGGCCTT CCCCACCCAGCAGGAAGCTCTGGGCACGCCACCCCCAAGTGCCCGATGTCTCTTGCCTGGTTCTTGCTACGG GACTCAACGATGGGCAGATCAAGATCTGGGAGGTGCAGACAGGGCTCCTGCTTTTGAATCTTTCCGGCCACCAAG ATGTCGTGAGAGATCTGAGCTTCACACCCAGTGGCAGTTTGATTTTGGTCTCCGCGTCACGGGATAAGACTCTTC GCATCTGGGACCTGAATAAACACGGTAAACAGATTCAAGTGTTATCGGGCCACCTGCAGTGGGTTTACTGCTGTT CCATCTCCCCAGACTGCAGCATGCTGTGCTCTGCAGCTGGAGAAGTCGGTCTTTCTATGGAGCATGAGGTCCT ACACGTTAATTCGGAAGCTAGAGGGCCATCAAAGCAGTGTTGTCTCTTGTGACTTCTCCCCCGACTCTGCCCTGC TTGTCACGGCTTCTTACGATACCAATGTGATTATGTGGGACCCCTACACCGGCGAAAGGCTGAGGTCACTCCACC ACACCCAGGTTGACCCCGCCATGGATGACAGTGACGTCCACATTAGCTCACTGAGATCTGTGTGCTTCTCCAG AAGGCTTGTACCTTGCCACGGTGGCAGATGACAGACTCCTCAGGATCTGGGCCCTGGAACTGAAAACTCCCATTG CATTTGCTCCTATGACCAATGGGCTTTGCTGCACATTTTTTCCACATGGTGGAGTCATTGCCACAGGGACAAGAG ATGGCCACGTCCAGTTCTGGACAGCTCCTAGGGTCCTGTCCTCACTGAAGCACTTATGCCGGAAAGCCCTTCGAA GTTTCCTAACAACTTACCAAGTCCTAGCACTGCCAATCCCCAAGAAAATGAAAGAGTTCCTCACATACAGGACTT TT**TAA**GCAACACCACATCTTGTGCTTCTTTGTAGCAGGGTAAATCGTCCTGTCAAAGGGAGTTGCTGGAATAATG GGCCAAACATCTGGTCTTGCATTGAAATAGCATTTCTTTGGGATTGTGAATAGAATGTAGCAAAACCAGATTCCA AACAGAGCCTTACGTTAAATTTTCAGTCCAGTTATGAACAGCAAGTGTTGAACTCTTTCTGCTTGTTTTGATTCA AAGTGCAGTTACTGATGTTTTTGATTATGCAACTAAGTAGGCCTCCAGAGCCTCTCTAGTGGCAGAGCAGCTC ACACTCCCTCCGCTGGGAACGATGGCTTCTGCCTAGTACCTATCCTTGTGTTTCTGATGCAGTGGTAGCATTGGT TCAAGTTCTCTCCTGCTGTGGTCAGAGTTGCTTCGATGTTGGCCAAGTGCTTTTCTTCTTGGGCTCCCTTCTGAC CTGCAGGACAGTTTTCCTGGAGCCATTTGGTATGAGGTATTAATTTAGCTTAACTAAATTACAGGGGACTCAGAG TGACAAATTTGTATGTCAGATTATACAAGGATGTATTCTTAAACCGCATGACTATTCAGATGGCTACTGAGTTAT CAGTGGCCATTTATTAGCATCATATTTATTTGTATTTTCTCAACAGATGTTAAGGTACAACTGTGTTTTTCTCGA TTATCTAAAAACCATAGTACTTAAATTGAACAGTTGCAAAGATGTCTTAATTGTGTAAAGAATTGGTGTAGTCAT GACTTTAGCTGATACTCTTATGTACGAGATCTGTCTCTGCTGTTTAACTTCATTGGATTAATCAGCTGGTTTCAA CTCTACTGCGAAACAAAAATAGCTCCTTAAAAGTACTGTTCTCCTTCAGTGGCATGTAGTTATCTAATCAAGACA CCTCATTCAAACAAACCTGCCTTAGGAAAATTTAATATATTTTAAATTATTTTAAAAGAAATACAACATCTTAT TCTTTAGCTTTCTTAATCGGTGCTTTATGGAGGCCAGTGTAACGTTACATGACTCGTTGAGAAAGTTGAGGAATT 

## 398/2825 FIGURE 363

MEAGEEPLLLAELKPGRPHQFDWKSSCETWSVAFSPDGSWFAWSQGHCIVKLIPWPLEEQFIPKGFEAKSRSSKN ETKGRGSPKEKTLDCGQIVWGLAFSPWPSPPSRKLWARHHPQVPDVSCLVLATGLNDGQIKIWEVQTGLLLLNLS GHQDVVRDLSFTPSGSLILVSASRDKTLRIWDLNKHGKQIQVLSGHLQWVYCCSISPDCSMLCSAAGEKSVFLWS MRSYTLIRKLEGHQSSVVSCDFSPDSALLVTASYDTNVIMWDPYTGERLRSLHHTQVDPAMDDSDVHISSLRSVC FSPEGLYLATVADDRLLRIWALELKTPIAFAPMTNGLCCTFFPHGGVIATGTRDGHVQFWTAPRVLSSLKHLCRK ALRSFLTTYQVLALPIPKKMKEFLTYRTF

### 399/2825 FIGURE 364

ATATAACCGCGTGGCCCGCGCGCGCTTCCCTCCCGGCGCGCTCACCGGCGCGCGTCTATGGCTGCGACTTCTCT  ${f AATG}$ TCTGCTTTGGCTGCCCGGCTGCAGCCCGCGCACAGCTGCTCCCTTCGCCCTTTCCACCTCGC GGCAGTTCGAAATGAAGCTGTTGTCATTTCTGGAAGGAAACTGGCCCAGCAGATCAAGCAGGAAGTGCGGCAGGA GGTAGAAGAGTGGGTGGCCTCAGGCAACAACGGCCACACCTGAGTGTGATCCTGGTTGGCGAGAATCCTGCAAG TCACTCCTATGTCCTCAACAAAACCAGGGCAGCTGCAGTTGTGGGAATCAACAGTGAGACAATTATGAAACCAGC TTCAATTTCAGAGGAAGAATTGTTGAATTTAATCAATAAACTGAATAATGATGATGATGATGGCCTCCTTGT TCAGTTGCCTCTTCCAGAGCATATTGATGAGAGAGGATCTGCAATGCTGTTTCTCCAGACAAGGATGTTGATGG CTTTCATGTAATTAATGTAGGACGAATGTGTTTGGATCAGTATTCCATGTTACCGGCTACTCCATGGGGTGTGTG GGAAATAATCAAGCGAACTGGCATTCCAACCCTAGGGAAGAATGTGGTTGTGGCTGGAAGGTCAAAAAACGTTGG AATGCCCATTGCAATGTTACTGCACACAGATGGGGCGCATGAACGTCCCGGAGGTGATGCCACTGTTACAATATC TCATCGATATACTCCCAAAGAGCAGTTGAAGAAACATACAATTCTTGCAGATATTGTAATATCTGCTGCAGGTAT TCCTGTAACTGCCAAACCCAAGTTGGTTGGAGATGTGGATTTTGAAGGAGTCAGACAAAAAGCTGGGTATATCAC TCCAGTTCCTGGAGGTGTTGGCCCCATGACAGTGGCAATGCTAATGAAGAATACCATTATTGCTGCAAAAAAGGT GCTGAGGCTTGAAGAGCGAGAAGTGCTGAAGTCTAAAGAGCTTGGGGTAGCCACTAAT**TAA**CTACTGTGTCTTCT CAGTACCTCACCAGGGAGCATTCCAGTATCATGCAGGGTCCTGTGATCTAGCCAGGAGCAGCCATTAACCTAGTG ATTAATATGGGAGACATTACCATATGGAGGATGCTTCACTTTGTCAAGCACCTCAGTTACACATTCGCCTT TTCTAGGATTGCATTTCCCAAGTGCTATTGCAATAACAGTTGATACTCATTTTAGGTACCAGACCTTTTGAGTTC AACTGATCAAACCAAAGGAAAAGTGTTGCTAGAGAAAATTGGGGAAAAAGGTGAAAAAAATGGTAGTAATT GAGCAGAAAAAATTAATTTATATGTATTGATTGGCAACCAGATTTATCTAAGTAGAACTGAATTGGCTAGGA AAAAAGAAAACTGCATGTTAATCATTTTCCTAAGCTGTCCTTTTGAGGCTTAGTCAGTTTATTGGGAAAATGTT AGGACTGAAAGGATTCTTTTCTACATTATACATGTGTGTTGTCATATTTTGGCTTTTTGCTATATACTTTAACTTCA AATCATTTGGGCAGCTTGGGTAAGTACGCAACTTACTTTTCCACCAAAGAACTGTCAGCAGCTGCCTGTTTTCT GTGATGTATCCTGTTGACTTTTCCAGAAATTTTTTAAGAGTTTGAGTTACTATTGAATTTAATCAGACTTT 

## 400/2825 FIGURE 365

MSALAARLLQPAHSCSLRLRPFHLAAVRNEAVVISGRKLAQQIKQEVRQEVEEWVASGNKRPHLSVILVGENPAS
HSYVLNKTRAAAVVGINSETIMKPASISEEELLNLINKLNNDDNVDGLLVQLPLPEHIDERRICNAVSPDKDVDG
FHVINVGRMCLDQYSMLPATPWGVWEIIKRTGIPTLGKNVVVAGRSKNVGMPIAMLLHTDGAHERPGGDATVTIS
HRYTPKEQLKKHTILADIVISAAGIPNLITADMIKEGAAVIDVGINRVHDPVTAKPKLVGDVDFEGVRQKAGYIT
PVPGGVGPMTVAMLMKNTIIAAKKVLRLEEREVLKSKELGVATN

# FIGURE 366

GGTAGTTGGTGGGCACTGGGTTAGAGGTATCACGTGGGGGCACTTTCGTCTTAGCTTTTGGACAAGACGCAG GGCGAACTGAGGCGAGTGAAGTGGACTCTGAGGGCTACCGCTACCGCCACTGCTGCGGCAGGGGCGTGGAGGGCA GAGGGCCGCGGAGGCCGCAGTTGCAAACATCGCTCAGAGCAGAGACGCGGAAACCCGTTCGCCGAGCCCAGCGA GCTTGACAACCCCTTTCAGGACCCAGCTGTGATCCAGCACCGACCCAGCCGGCAGTATGCCACGCTTGACGTCTA CAACCCTTTTGAGACCCGGGAGCCACCACCACCTTTGAGCCTCCAGCCCTTGCCCCATTGCCTCCACCTCAGC TCCCTCCTTGCAGCCCTCGAGAAAGCTCAGCCCCACAGAACCTAAGAACTATGGCTCATACAGCACTCAGGCCTC AGCTGCAGCAGCCACAGCTGAGCTGCTGAAGAAACAGGAGGAGCTCAACCGGAAGGCAGAGGAGTTGGACCGAAG GGAGCGAGAGCTGCAGCATGCTGCCCTGGGGGGCACAGCTACTCGACAGAACAATTGGCCCCCTCTACCTTCTTT TTGTCCAGTTCAGCCCTGCTTTTTCCAGGACATCTCCATGGAGATCCCCCAAGAATTTCAGAAGACTGTATCCAC GGAAACCAACAATGGCGCAGGCTTTGGGCTTTCTATCCTCTGGGTCCTCCTTTTCACTCCCTGCTCCTTTGTCTG  $\tt CGTCCAGGATGTGCTCTTTGTCCTCCAGGCCATTGGTATCCCAGGTTGGGGATTCAGTGGCTGGATCTCTGCTCT$ GGTGGTGCCGAAGGGCAACACAGCAGTATCCGTGCTCATGCTGGTCGCCCTGCTCTTCACTGGCATTGCTGT GCTAGGAATTGTCATGCTGAAACGGATCCACTCCTTATACCGCCGCACAGGTGCCAGCTTTCAGAAGGCCCAGCA AGAATTTGCTGCTGGTGTCTTCTCCAACCCTGCGGTGCGAACCGCAGCTGCCAATGCAGCCGCTGGGGCTGCTGA AAATGCCTTCCGGGCCCCG**TGA**CCCCTGACTGGGATGCCCTGGCCCTGCTACTTGAGGGAGCTGACTTAGCTCCC GTCCCTAAGGTCTCTGGGACTTGGAGAGACATCACTAACTGATGGCTCCTCCGTAGTGCTCCCAATCCTATGGCC ATGACTGCTGAACCTGACAGGCGTGTGGGGAGTTCACTGTGACCTAGTCCCCCCATCAGGCCACACTGCTGCCAC 

## 402/2825 FIGURE 367

MAQSRDGGNPFAEPSELDNPFQDPAVIQHRPSRQYATLDVYNPFETREPPPAYEPPAPAPLPPPSAPSLQPSRKL SPTEPKNYGSYSTQASAAAATAELLKKQEELNRKAEELDRRERELQHAALGGTATRQNNWPPLPSFCPVQPCFFQ DISMEIPQEFQKTVSTMYYLWMCSTLALLLNFLACLASFCVETNNGAGFGLSILWVLLFTPCSFVCWYRPMYKAF RSDSSFNFFVFFIFFVQDVLFVLQAIGIPGWGFSGWISALVVPKGNTAVSVLMLLVALLFTGIAVLGIVMLKRI HSLYRRTGASFQKAQQEFAAGVFSNPAVRTAAANAAAGAAENAFRAP

# FIGURE 368

CCACGTGGTGTTACATATATTATTTTCATACATGGTTTCTGGCTCATAAGTTCCTTAGCCCTTGCTATAGTCT TTTGTGTTCGGTCTTAAGGGCAGGACTGTACTCTTCCCTCACCTTTCTAATTGTGCATCTTAAGACCTTCCCCAG AGAGGGTGCCCTGTAGTTGTGGGAAGGAATGCTGGCATCATGAAGCTTCCATAAAAACCCGAGAAACGAGCT  ${\tt TCTGGATAGCTGGACACATGGAGGTCCTGGAGGGTGGAGCCCAGGGAGGCATGGAAGCTCCACAGCCCTTCCCCC}$ ATACCTTACCCTATTTCCTCTGTATCCTTTGTAATATCCTTTATGATAAACCAGCAAATGTGTGTAAATGTTTCC CTAAGGTCTGTGGCCACTCCAGCAAATTAATTGAACCTAAAGAGGGGGTCGTGGGAACCCCAACTTGAAGCCAGT CAGTCAGAAGTTCTGGATGTCCAGACTTCAGACTGGTGTCTGAAAGGGTGGAGGCAGTCTTGGGGACCGAGCCCC CAATCTATGGGATCTGACACTATCTCCAGTAGTGTTGGAATTGAGTCACCAGCGTGTCCACTGGTTAGTGTGTA GAAACTCCCTACCATTGGTCACAGAAGTCTTCTTCTGTGTTGATAGTTGTAGTGTGACAGCAGAGGAAAAACAAA GTCAGAAAGAGTTTTCCCGAACACCCCAATTTCTCCATTTTACTATCCATTTCCACAAACACTGACTACAATAG TTGTAGGTCACCACCTAGAGGGGAGGAAGACCTCGCTTTGGAGAGTGGGAATAAAACGCTCGTGGAAAAGGGTAC ACGCTTTTCTGGGAAAGTGAGGCCACC**ATG**GCTCTGGAGAAGTCTCTTGTCCGGCTCCTTCTGCTTGTCCTGATA CTGCTGGTGCTGGGCTCGGCCTTCCCTGGGCAAGGAATCCCGGGCCAAGAAATTCCAGCGGCAGCATATG GACTCAGACAGTTCCCCCAGCAGCAGCTCCACCTACTGTAACCAAATGATGAGGCGCCGGAATATGACACAGGGG CGGTGCAAACCAGTGAACACCTTTGTGCACGAGCCCCTGGTAGATGTCCAGAATGTCTGTTTCCAGGAAAAGGTC ACCTGCAAGAACGGGCAGGCAACTGCTACAAGAGCAACTCCAGCATGCACATCACAGACTGCCGCCTGACAAAC GGCTCCAGGTACCCCAACTGTGCATACCGGACCAGCCCGAAGGAGACACATCATTGTGGCCTGTGAAGGGAGC CTCCTATTCAACACACACATGCTTCCCTTTCCTGAGCCGGAATTC

## 404/2825 FIGURE 369

 ${\tt MALEKSLVRLLLVLILLVLGWVQPSLGKESRAKKFQRQHMDSDSSPSSSSTYCNQMMRRRNMTQGRCKPVNTFV}\\ {\tt HEPLVDVQNVCFQEKVTCKNGQGNCYKSNSSMHITDCRLTNGSRYPNCAYRTSPKERHIIVACEGSPYVPVHFDASVEDST}\\$ 

### 405/2825 FIGURE 370A

CTTCTCTCCCAGGGCGGTGCGACCCGGAGCTCCAGCGCCCGAGTCTCCACTTCGTTTGCTGAAACTTGCTTTCTA CCAGCTAAGAACCATGCTGCGAGTGATTGTGGAATCTGCCAGCAATATCCCTAAAACGAAATTTGGCAAGCCGGA GAATGAGATTTTGGAGTTTGACTTGAGGGGTATACCACTGGACTTTTCATCTTCCCTTGGGATTATTGTGAAAGA TTTTGAGACAATTGGACAAATTAATTGGCACGGCGACTGTAGCCCTGAAGGACCTGACTGGTGACCAGAG CAGATCCCTGCCGTACAAGCTGATCTCCCTGCTAAATGAAAAAGGGCCAAGATACTGGGGCCCACTAGACTTGGT GATCGGCTATGATCCGCCTTCTGCTCCACATCCAAATGACCTGAGCGGGCCCAGCGTGCCAGGCATGGGAGGAGA TGGGGAAGAAGATGAAGGTGATGAAGACAGGTTGGACAATGCAGTCAGGGGCCCTGGGCCCAAGGGGCCCAGTTGG GACGGTGTCGGAAGCTCAGCTTGCTCGGAGGCTCACCAAAGTAAAGAACAGCCGGCGGATGCTGTCAAATAAGCC ACAGGACTTCCAGATCCGCGTCCGAGTGATTGAGGGCCGACAGTTAAGTGGTAACAACATAAGGCCTGTGGTCAA AGTTCACGTCTGTGGCCAGACACCCGAACAAGAATCAAGAGAGGAAACAACCCTTTTTTTGATGAGTTGTTTTT CTACAATGTCAACATGACCCCTTCTGAATTGATGGATGAGATCATCAGCATCCGGGTTTATAATTCTCACTCTC GAGAAAGTGGCTTCTTCTCAATGACCCGGAAGATACCAGTTCAGGTTCTAAAGGTTATATGAAAGTCAGCATGTT TGTCCTGGGAACCGGAGATGAGCCTCCTCCTGAGAGACGAGATCGTGATAATGACAGTGATGATGTGGAGAGTAA TTTGTTACTCCCTGCTGGCATTGCCCTCCGGTGGGTGACCTTCTTGCTGAAAATCTACCGAGCTGAGGACATCCC CCAGATGGATGATGCCTTCTCACAGACAGTAAAGGAAATATTTGGAGGCAATGCAGATAAGAAAAATCTCGTGGA TCCTTTTGTAGAAGTTTCCTTTGCTGGAAAAAAGGTTTGTACAAACATAATTGAGAAAAATGCAAACCCAGAGTG GAATCAGGTCGTCAATCTTCAGATCAAGTTTCCTTCAGTGTGTGAAAAAATAAAACTAACAATATATGACTGGGA CCGTCTTACTAAAAATGATGTAGTTGGAACAACATATCTACACCTCTCTAAAATTGCTGCCTCTGGTGGGGAAGT GGAAGATTTCTCATCTTCGGGAACTGGGGCTGCATCATATACAGTAAACACAGGAGAAACAGAGGTAGGCTTTGT TCCAACGTTTGGACCTTGTTACCTGAATCTTTATGGAAGCCCCAGGGAGTACACGGGATTCCCAGACCCCTATGA TGAGCTGAATACTGGAAAGGGGGAAGGAGTTGCCTACAGAGGCAGGATCTTGGTTGAATTAGCCACTTTTCTTGA GAAGACACCACCAGATAAAAAGCTTGAGCCCATTTCAAATGATGACCTGCTGGTTGTTGAGAAATACCAGCGAAG GCGGAAGTACAGCCTGTCTGCCGTGTTTCATTCAGCCACCATGTTGCAAGATGTTGGTGAGGCCATTCAGTTTGA AGTCAGCATTGGGAACTATGGCAACAAGTTTGACACCACCTGTAAGCCTTTGGCATCAACAACTCAGTACAGCCG TGCTGTATTTGATGGCAACTACTATTATTACTTGCCTTGGGCCCACACCAAGCCAGTTGTTACCCTGACTTCATA CTGGGAGGATATTAGTCATCGCCTGGATGCGGTGAACACTCTCCTAGCTATGGCAGAACGGCTGCAAACAATAT TGAAGTTATAGAAGACACGAGATACACGTTGCCTCTCACAGAAGGAAAAGCCAACGTCACAGTTCTCGATACTCA GATCCGAAAGCTGCGGTCCAGGTCTCTCTCCCAAATACATGAGGCGGCTGTGAGGATGAGGTCGGAAGCCACAGA CTTGTACTCCACCAGTGGTGAGAATGCATCTGGGAAAATACTGTGGGAAAACCCAAACCATCTTTCTGAAGTATCC ACAGGAGAAAAACAACGGGCCAAAGGTGCCTGTGGAGTTGCGAGTGAACATCTGGCTAGGCTTAAGTGCTGTGGA GAAGAAGTTTAACAGCTTCGCAGAAGGAACTTTCACCGTCTTTGCTGAAAATGTATGAAAATCAAGCTCTCATGTT TGGAAAATGGGGTACTTCTGGATTAGTAGGACGTCATAAGTTTTCTGATGTCACAGGAAAAATAAAACTCAAGAG GGAATTTTTTCTGCCTCCAAAAGGCTGGGAATGGGAAGGAGTGGATAGTTGATCCTGAAAGAAGCTTGCTGAC TGAGGCAGATGCAGGTCACACGGAGTTCACTGATGAAGTCTATCAGAACGAGAGCCGCTACCCCGGGGGCGACTG GAAGCCGGCCGAGGACACCTACACGGATGCGAACGGCGATAAAGCAGCATCACCCAGCGAGTTGACTTGTCCTCC AGGTTGGGAATGGGAAGATGATGCATGGTCTTATGACATAAATCGAGCGGTGGATGAGAAAGGCTGGGAATATGG AATCACCATTCCTCCTGATCATAAGCCCAAATCCTGGGTTGCAGCAGAAAAATGTACCACACTCATAGACGGCG AAGGCTGGTCCGAAAACGCAAGAAAGATTTAACACAGACTGCTTCAAGCACCGCAAGGGCCATGGAGGAATTGCA AGACCAAGAGGGCTGGGAATATGCTTCTCTAATTGGCTGGAAATTTCACTGGAAACAACGTAGTTCAGATACCTT CCGCCGCAGACGCTGGAGGAGAAAAATGGCTCCTTCAGAAACACATGGTGCAGCTGCCATCTTTAAACTTGAAGG TGCCCTTGGGGCAGACACTACCGAAGATGGGGATGAGAAGACCTGGAGAAACAGAAGCACAGTGCCACCACTGT GTTCGGAGCAAACACCCCCATTGTTTCCTGCAATTTTGACAGAGTCTACATCTACCATCTGCGCTGCTATGTCTA TCAAGCCAGAAACCTCTTGGCTTTAGATAAGGATAGCTTTTCAGATCCATATGCTCATATCTGTTTCCTCCATCG

### 406/2825 FIGURE 370B

GAGCAAAACCACTGAGATCATCCATTCAACCCTGAATCCCACGTGGGACCAAACAATTATATTCGATGAAGTTGA AATCTATGGGGAACCCCAAACAGTTCTACAGAATCCACCCAAAGTTATCATGGAACTTTTTGACAATGACCAAGT CAAACTTCTCTGGCACCCAGTAATGAATGGAGACAAAGCCTGCGGGGGATGTTCTTGTAACTGCAGAGCTGATTCT GAGGGGCAAGGATGGCTCCAACCTTCCCATTCTTCCCCCTCAAAGGGCGCCAAATCTATACATGGTCCCCCAGGG GATCAGGCCTGTGGTCCAGCTCACTGCCATTGAGAATTCTAGCTTGGGGCTTAAGAAATATGAAAAACTTCCAGAT GGCTTCTATCACATCCCCCAGTCTTGTTGTGGAGTGTGGAGGAGAAAGGGTGGAATCGGTGGTGATCAAAAACCT TAAGAAGACACCCAACTTTCCAAGTTCTGTTCTCTTCATGAAAGTGTTCTTGCCCAAGGAGGAATTGTACATGCC CCTGGACCGCTTTCGCTGTGACCCTTATGCAGGGAAAGAGGACATCGTCCCACAGCTCAAAGCCTCCCTTCTGTC TGCCCCACCATGCCGGGACATCGTTATCGAAATGGAAGACACCAAACCATTACTGGCTTCTAAGCTGACAGAAAA GGAGGAAGAATCGTGGACTGGTGGAGTAAATTTTATGCTTCCTCAGGGGAACATGAAAAATGCGGACAGTATAT TCAGAAAGGCTATTCCAAGCTCAAGATATATATTGTGAACTAGAAAATGTAGCAGAATTTGAGGGCCTGACAGA CTTCTCAGATACGTTCAAGTTGTACCGAGGCAAGTCGGATGAAAATGAAGATCCTTCTGTGGTTGGAGAGTTTAA GGGCTCCTTTCGGATCTACCCTCTGCCGGATGACCCCAGCGTGCCAGCCCCTCCCAGACAGTTTCGGGAATTACC TGACAGCGTCCCACAGGAATGCACGGTTAGGATTTACATTGTTCGAGGCTTAGAGCTCCAGCCCCAGGACAACAA TGGCCTGTGTGACCCTTACATAAAAATAACACTGGGCAAAAAAGTCATTGAAGACCGAGATCACTACATTCCCAA CACTCTCAACCCAGTCTTTGGCAGGATGTACGAACTGAGCTGCTACTTACCTCAAGAAAAAGACCTGAAAATTTC TGTCTATGATTATGACACCTTTACCCGGGATGAAAAAGTAGGAGAAACAATTATTGATCTGGAAAACCGATTCCT TTCCCGCTTTGGGTCCCACTGCGGCATACCAGAGGAGTACTGTGTTTCTGGAGTCAATACCTGGCGAGATCAACT GAGACCAACACAGCTGCTTCAAAATGTCGCCAGATTCAAAGGCTTCCCACAACCCATCCTTTCCGAAGATGGGAG TAGAATCAGATATGGAGGACGAGACTACAGCTTGGATGAATTTGAAGCCAACAAAATCCTGCACCAGCACCTCGG GGCCCCTGAAGAGCGGCTTGCTCTTCACATCCTCAGGACTCAGGGGCTGGTCCCTGAGCACGTGGAAACAAGGAC GGGGCCACCAGGCCCTCCTTTCAACATCACCCCCGGAAAGCCAAGAAATACTACCTGCGTGTGATCATCTGGAA CACCAAGGACGTTATCTTGGACGAGAAAAGCATCACAGGAGAGAAATGAGTGACATCTACGTCAAAGGCTGGAT TCCTGGCAATGAAGAAACAAACAGAAACAGATGTCCATTACAGATCTTTGGATGGTGAAGGGAATTTTAACTG GCGATTTGTTTTCCCGTTTGACTACCTTCCAGCCGAACAACTCTGTATCGTTGCGAAAAAAAGAGCATTTCTGGAG TATTGACCAAACGGAATTTCGAATCCCACCCAGGCTGATCATTCAGATATGGGACAATGACAAGTTTTCTCTGGA TGACTACTTGGGTTTCCTAGAACTTGACTTGCGTCACACGATCATTCCTGCAAAATCACCAGAGAAATGCAGGTT GAAAGGATGGTGGCCATGCTACGCAGAGAAAGATGGCGCCCCGCGTAATGGCTGGGAAAGTGGAGATGACATTGGA AATCCTCAACGAGAAGGAGGCCGACGAGAGGCCAGCCGGGAAGGGGCGGACGAACCCAACATGAACCCCAAGCT GGACTTACCAAATCGACCAGAAACCTCCTTCCTCTGGTTCACCAACCCATGCAAGACCATGAAGTTCATCGTGTG GCGCCGCTTTAAGTGGGTCATCATCGGCTTGCTGTTCCTGCTTATCCTGCTGCTCTTCGTGGCCGTGCTCCTCTA CTCTTTGCCGAACTATTTGTCAATGAAGATTGTAAAGCCAAATGTGTAACAAAGGCAAAGGCTTCATTTCAAGAG TCATCCAGCAATGAGAGAATCCTGCCTCTGTAGACCAACATCCAGTGTGATTTTGTGTCTGAGACCACACCCCAG TAGCAGGTTACGCCATGTCACCGAGCCCCATTGATTCCCAGAGGGTCTTAGTCCTGGAAAGTCAGGCCAACAAGC AACGTTTGCATCATGTTATCTCTTAAGTATTAAAAGTTTTATTTTCTAAAGTTTAAATCATGTTTTTCAAAATAT TTTTCAAGGTGGCTGGTTCCATTTAAAAATCATCTTTTTATATGTGTCTTCGGTTCTAGACTTCAGCTTTTGGAA ATTGCTAAATAGAATTCAAAAATCTCTGCATCCTGAGGTGATATACTTCATATTTGTAATCAACTGAAAGAGCTG AAAA

### 407/2825 FIGURE 371

MLRVIVESASNIPKTKFGKPDPIVSVIFKDEKKKTKKVDNELNPVWNEILEFDLRGIPLDFSSSLGIIVKDFETI GQNKLIGTATVALKDLTGDQSRSLPYKLISLLNEKGQDTGATIDLVIGYDPPSAPHPNDLSGPSVPGMGGDGEED EGDEDRLDNAVRGPGPKGPVGTVSEAQLARRLTKVKNSRRMLSNKPQDFQIRVRVIEGRQLSGNNIRPVVKVHVC GQTHRTRIKRGNNPFFDELFFYNVNMTPSELMDEIISIRVYNSHSLRADCLMGEFKIDVGFVYDEPGHAVMRKWL LLNDPEDTSSGSKGYMKVSMFVLGTGDEPPPERRDRDNDSDDVESNLLLPAGIALRWVTFLLKIYRAEDIPOMDD AFSQTVKEIFGGNADKKNLVDPFVEVSFAGKKVCTNIIEKNANPEWNQVVNLQIKFPSVCEKIKLTIYDWDRLTK NDVVGTTYLHLSKIAASGGEVEDFSSSGTGAASYTVNTGETEVGFVPTFGPCYLNLYGSPREYTGFPDPYDELNT GKGEGVAYRGRILVELATFLEKTPPDKKLEPISNDDLLVVEKYQRRRKYSLSAVFHSATMLQDVGEAIOFEVSIG NYGNKFDTTCKPLASTTQYSRAVFDGNYYYYLPWAHTKPVVTLTSYWEDISHRLDAVNTLLAMAERLQTNIEALK SGIQGKIPANQLAELWLKLIDEVIEDTRYTLPLTEGKANVTVLDTQIRKLRSRSLSQIHEAAVRMRSEATDVKST LAEIEDWLDKLMQLTEEPQNSMPDIIIWMIRGEKRLAYARIPAHQVLYSTSGENASGKYCGKTQTIFLKYPQEKN NGPKVPVELRVNIWLGLSAVEKKFNSFAEGTFTVFAEMYENQALMFGKWGTSGLVGRHKFSDVTGKIKLKREFFL PPKGWEWEGEWIVDPERSLLTEADAGHTEFTDEVYQNESRYPGGDWKPAEDTYTDANGDKAASPSELTCPPGWEW EDDAWSYDINRAVDEKGWEYGITIPPDHKPKSWVAAEKMYHTHRRRRLVRKRKKDLTQTASSTARAMEELQDQEG WEYASLIGWKFHWKQRSSDTFRRRRWRRKMAPSETHGAAAIFKLEGALGADTTEDGDEKSLEKQKHSATTVFGAN TPIVSCNFDRVYIYHLRCYVYQARNLLALDKDSFSDPYAHICFLHRSKTTEIIHSTLNPTWDQTIIFDEVEIYGE PQTVLQNPPKVIMELFDNDQVGKDEFLGRSIFSPVVKLNSEMDITPKLLWHPVMNGDKACGDVLVTAELILRGKD GSNLPILPPQRAPNLYMVPQGIRPVVQLTAIEILAWGLRNMKNFQMASITSPSLVVECGGERVESVVIKNLKKTP NFPSSVLFMKVFLPKEELYMPPLVIKVIDHRQFGRKPVVGQCTIERLDRFRCDPYAGKEDIVPQLKASLLSAPPC RDIVIEMEDTKPLLASKLTEKEEEIVDWWSKFYASSGEHEKCGQYIQKGYSKLKIYNCELENVAEFEGLTDFSDT FKLYRGKSDENEDPSVVGEFKGSFRIYPLPDDPSVPAPPRQFRELPDSVPQECTVRIYIVRGLELOPODNNGLCD PYIKITLGKKVIEDRDHYIPNTLNPVFGRMYELSCYLPOEKDLKISVYDYDTFTRDEKVGETIIDLENRFLSRFG SHCGIPEEYCVSGVNTWRDQLRPTQLLQNVARFKGFPQPILSEDGSRIRYGGRDYSLDEFEANKILHOHLGAPEE RLALHILRTQGLVPEHVETRTLHSTFQPNISQGKLQMWVDVFPKSLGPPGPPFNITPRKAKKYYLRVIIWNTKDV ILDEKSITGEEMSDIYVKGWIPGNEENKQKTDVHYRSLDGEGNFNWRFVFPFDYLPAEOLCIVAKKEHFWSIDOT EFRIPPRLIIQIWDNDKFSLDDYLGFLELDLRHTIIPAKSPEKCRLDMIPDLKAMNPLKAKTASLFEQKSMKGWW PCYAEKDGARVMAGKVEMTLEILNEKEADERPAGKGRDEPNMNPKLDLPN

### 408/2825 FIGURE 372

GGGCTGCGCTGTCCAGCTGTGGCTATGGCCCCAGCCCCGAGATGAGGAGGAGAGAACTAGGGGCCCGCAGGCCT GCCAGGGAAAACCGAGACACCATCACCATCACCAGTCACCAGCCTCAGGACAGATACAAAGCTGTCTGGCT TATCTTCTTCATGCTGGGTCTGGGAACGCTGCTCCCGTGGAATTTTTTCATGACGGCCACTCAGTATTTCACAAA CCGCCTGGACATGTCCCAGAATGTGTCCTTGGTCACTGCTGAACTGAGCAAGGACGCCCAGGCGTCAGCCGCCC TGCAGCACCCTTGCCTGAGCGGAACTCTCTCAGTGCCATCTTCAACAATGTCATGACCCTATGTGCCATGCTGCC CCTGCTGTTATTCACCTACCTCAACTCCTTCCTGCATCAGAGGATCCCCCAGTCCGTACGGATCCTGGGCAGCCT GGTGGCCATCCTGCTGGTGTTTCTGATCACTGCCATCCTGGTGAAGGTGCAGCTGGATGCTCTGCCCTTCTTTGT CCTTCTGCCTGCCAGCTACACGGCCCCCATCATGAGTGGCCAGGGCCTAGCAGGCTTCTTTGCCTCCGTGGCCAT GATCTGCGCTATTGCCAGTGGCTCGGAACTATCAGAAAGTGCCTTCGGCTACTTTATCACAGCCTGTGCTGTTAT CATTTTGACCATCATCTGTTACCTGGGCCTGCCCCGCCTGGAATTCTACCGCTACTACCAGCAGCTCAAGCTTGA AGGACCCGGGGAGCCAGGAGCCAAGTTGGACCTCATTAGCAAAGGAGGAGCCCAAGAGCAGGCAAAGAGGAATC TGGAGTTTCAGTCTCCAACTCTCAGCCCACCAATGAAAGCCACTCTATCAAAGCCATCCTGAAAAATATCTCAGT CCTGGCTTTCTCTGTCTGCTTCATCTTCACTATCACCATTGGGATGTTTCCAGCCGTGACTGTTGAGGTCAAGTC CAGCATCGCAGGCAGCACCTGGGAACGTTACTTCATTCCTGTGTCCTGTTTCTTGACTTTCAATATCTTTGA GGCCCGGCTGGTTTTGTGCCACTGCTGCTGCTGCAACATTAAGCCCCGCCGCTACCTGACTGTGGTCTTCGA GCACGATGCCTGGTTCATCTTCTTCATGGCTGCCTTTGCCTTCTCCAACGGCTACCTCGCCAGCCTCTGCATGTG CTTCGGGCCCAAGAAAGTGAAGCCAGCTGAGGCAGAGACCGCAGGAGCCATCATGGCCTTCTTCCTGTGTCTGGG  $\tt CCTCCCTGCCTGCCTCCTGCCCCTTCCTTCTGCCAGGGGTGATCCTGAGTGGTCTGGCGGTTTTTTCTTCT$ GTGGGGACATTGTGGGTTTGGGGCTCAGAGTCGAGGGACGGGGTGTAGCCTCGGCATTTGCTTGAGTTTCTCCAC TCTTGGCTCTGACTGATCCCTGCTTGTGCAGGCCAGTGGAGGCTCTTGGGCTTTGGAGAACACGTGTGTCTCTGTG CGTTGTATGGTTTGACCTGATATACTCCATTCTCCCCTGCGCCTCCTCTGTGTTTTTTTCCATGTCCCCCTCC CAACTCCCCATGCCCAGTTTTTACCCATCATGCACCCTGTACAGTTGCCACGTTACTGCCTTTTTTAAAAATATA 

## 409/2825 FIGURE 373

MTTSHQPQDRYKAVWLIFFMLGLGTLLPWNFFMTATQYFTNRLDMSQNVSLVTAELSKDAQASAAPAAPLPERNS LSAIFNNVMTLCAMLPLLLFTYLNSFLHQRIPQSVRILGSLVAILLVFLITAILVKVQLDALPFFVITMIKIVLI NSFGAILQGSLFGLAGLLPASYTAPIMSGQGLAGFFASVAMICAIASGSELSESAFGYFITACAVIILTIICYLG LPRLEFYRYYQQLKLEGPGEQETKLDLISKGEEPRAGKEESGVSVSNSQPTNESHSIKAILKNISVLAFSVCFIF TITIGMFPAVTVEVKSSIAGSSTWERYFIPVSCFLTFNIFDWLGRSLTAVFMWPGKDSRWLPSLVLARLVFVPLL LLCNIKPRRYLTVVFEHDAWFIFFMAAFAFSNGYLASLCMCFGPKKVKPAEAETAGAIMAFFLCLGLALGAVFSF LFRAIV

# FIGURE 374

CCTGGGCCGGCCGGGTGATGAGCCGGGAGCTCCCTGCTGCCGGTCATACCACAGCCTTCATCTGCGCCCTGGG CCGTCGCTCCGGCCGCCTCCCCTCGGGTCCCCGTGCGAGCCCACGCCGGCCCCGGTGCCCGCAGCCCT GCCACTGGACACAGGATAAGGCCCAGCGCACAGGCCCCCACGTGGACAGCATCGACCGCGGCACGCTCCCTCTGG  $\tt CTGTTGCCCTGCTGCCAGCCTGCAGCCTCAGCCCCACAAGTCTTGCAGAAACAGTCCATTGTGACCTTCAGC$ ATGCCATCCTTGAAGTCCATGTCCTCTTCCTGGAGTTCCCAACGGGCCCGTCACAGCTGGAGCTGACTCTCCAGG  ${\tt CATCCAAGCAAAATGGCACCTGGCCCCGAGAGGTGCTTCTGGTCCTCAGTGTAAACAGCAGTGTCTTCCTGCATC}$ TCCAGGCCTGGGAATCCCACTGCACTTGGCCTACAATTCCAGCCTGGTCACCTTCCAAGAGCCCCCGGGGGTCA ACACCACAGAGCTGCCATCCTTCCCCAAGACCCAGATCCTTGAGTGGGCAGCTGAGAGGGGCCCCATCACCTCTG TGGAAGCCAGCCAGGACATGGGCCGCACGCTCGAGTGGCGGCCGCGTACTCCAGCCTTGGTCCGGGGCTGCCACT TGGAAGGCGTGGCCGGCCACAAGGAGGCGCACATCCTGAGGGTCCTGCCGGGCCACTCGGCCGGGCCCCGGACGG TGTCCTGGCTCATCGACGCCAACCACACATGCAGATCTGGACCACTGGAGAATACTCCTTCAAGATCTTTCCAG AGAAAAACATTCGTGGCTTCAAGCTCCCAGACACACCTCAAGGCCTCCTGGGGGAGGCCCGGATGCTCAATGCCA GCATTGTGGCATCCTTCGTGGAGCTACCGCTGGCCAGCATTGTCTCACTTCATGCCTCCAGCTGCGGTGGTAGGC TGCAGACCTCACCGGACCGATCCAGACCACTCCTCCCAAGGACACTTGTAGCCCGGAGCTGCTCATGTCCTTGA TCACGGGCCTGACCTTCTGGGACCCCAGCTGTGAGGCAGAGGACAGGGGTGACAAGTTTGTCTTGCGCAGTGCTT ACTCCAGCTGTGGCATGCAGGTGTCAGCAAGTATGATCAGCAATGAGGCGGTGGTCAATATCCTGTCGAGCTCAT CACCACAGCGGAAAAAGGTGCACTGCCTCAACATGGACAGCCTCTCTTTCCAGCTGGGCCTCTACCTCAGCCCAC ACTTCCTCCAGGCCTCCAACACCATCGAGCCGGGGCAGCAGAGCTTTGTGCAGGTCAGAGTGTCCCCATCCGTCT GCCGGCCGCCCAAGGCCAACTGTGTGAGCCTGCTGTCCCCAAGCCCCGAGGGTGACCCGCGCTTCAGCTTCCTCC TCCACTTCTACACAGTACCCATACCCAAAACCGGCACCCTCAGCTGCACGGTAGCCCTGCGTCCCAAGACCGGGT CTCAAGACCAGGAAGTCCATAGGACTGTCTTCATGCGCTTGAACATCATCAGCCCTGACCTGTCTGGTTGCACAA GCAAAGGCCTCGTCCTGCCCGCCGTGCTGGGCATCACCTTTGGTGCCTTCCTCATCGGGGCCCTGCTCACTGCTG CACTCTGGTACATCTACTCGCACACGCGTGAGTACCCCAGGCCCCCACAG**TGA**GCATGCCGGGCCCCTCCATCCA CCCGGGGGAGCCCAGTGAAGCCTCTGAGGGATTGAGGGGCCCTGGCAGGACCCTGACCTCCGCCCCTGCCCCGC TCCCGCTCCCAGGTTCCCCCAGCAAGCGGGAGCCCGTGGTGGCGGTGGCTGCCCCGGCCTCCTCGGAGAGCAGCA GCACCAACCACAGCATCGGGAGCACCCAGAGCACCCCTGCTCCACCAGCAGCATGGCATAGCCCCGGCCCCCCG CGCTCGCCCAGCAGGAGAGACTGAGCAGCCGCCAGCTGGGAGCACTGGTGTGAACTCACCCTGGGAGCCAGTCCT AGCCACTGGCTTGGAACACCTTGGGGTCCCTCCACCCCACAGAACCTTCAACCCAGTGGGTCTGGGATATGGCTG  $\verb|CCCAGGAGACCACTTGCCACGCTGTTGTAAAAACCCAAGTCCCTGTCATTTGAACCTGGATCCAGCACTGG|$  $\tt CTGGGAAGAGAGGGCCCAGCCCAGAGCCACCTGGATCTATCCCTGCGCCTCCACACCTGAACTTGCCTAAC$ TAACTGGCAGGGGAGACAGGAGCCTAGCGGAGCCCAGCCTGGGAGCCCAGAGGGTGGCAAGAACAGTGGGCGTTG 

## 411/2825 FIGURE 375

MDRGTLPLAVALLLASCSLSPTSLAETVHCDLQPVGPERGEVTYTTSQVSKGCVAQAPNAILEVHVLFLEFPTGP
SQLELTLQASKQNGTWPREVLLVLSVNSSVFLHLQALGIPLHLAYNSSLVTFQEPPGVNTTELPSFPKTQILEWA
AERGPITSAAELNDPQSILLRLGQAQGSLSFCMLEASQDMGRTLEWRPRTPALVRGCHLEGVAGHKEAHILRVLP
GHSAGPRTVTVKVELSCAPGDLDAVLILQGPPYVSWLIDANHNMQIWTTGEYSFKIFPEKNIRGFKLPDTPQGLL
GEARMLNASIVASFVELPLASIVSLHASSCGGRLQTSPAPIQTTPPKDTCSPELLMSLIQTKCADDAMTLVLKKE
LVAHLKCTITGLTFWDPSCEAEDRGDKFVLRSAYSSCGMQVSASMISNEAVVNILSSSSPQRKKVHCLNMDSLSF
QLGLYLSPHFLQASNTIEPGQQSFVQVRVSPSVSEFLLQLDSCHLDLGPEGGTVELIQGRAAKGNCVSLLSPSPE
GDPRFSFLLHFYTVPIPKTGTLSCTVALRPKTGSQDQEVHRTVFMRLNIISPDLSGCTSKGLVLPAVLGITFGAF
LIGALLTAALWYIYSHTREYPRPPQ

# FIGURE 376

CTGGAGGAGGAAGAGGTGGATCCCCGGATCCAGGGAGAACTGGAGAAGTTAAATCAGTCCACGGATGATATC AACAGACGGGAGACTGAACTTGAGGATGCTCGTCAGAAGTTCCGCTCTGTTCTGGTTGAAGCAACGGTGAAACTG GATGAACTGGTGAAGAAATTGGCAAAGCTGTGGAAGACTCCAAGCCCTACTGGGAGGCACGGAGGGTGGCGAGG CAGGCTCAGCTGGAAGCTCAGAAAGCCACGCAGGACTTCCAGAGGGCCCACAGAGGTGCTCCGTGCCGCCAAGGAG ACCATCTCCCTGGCCGAGCAGCAGCTGCTGGAGGATGACAAGCGGCAGTTCGACTCCGCCTGGCAGGAGATGCTG GCCAGGTACAATGCCGCCATGGGCCGCATGCGACAGCTGGAGAAAACTCAAGAGGCCATCAACAAGTCCAAG CCTTATTTTGAACTCAAGGCAAAGTACTATGTGCAGCTCGAGCAACTGAAAAAGACTGTGGATGACCTGCAGGCC AAACTGACCCTGGCAAAAGGCGAGTACAAGATGGCCCTGAAGAACCTGGAGATGATCTCAGATGAGATCCACGAG TTTGTGTCTGAAGATGACTCGGAAACCCAGTCCGTGTCCAGCTTTAGTTCAGGACCAACAAGCCCGTCTGAGATG  $\verb|CCTGACCAGTTCCCTGCGGTTGTGAGGCCTGGCAGCCTGGATCTGCCCAGCCCTGTGTCCAGAGTTTGGG| \\$ ATGATGTTCCCAGTGTTGGGCCCTCGAAGTGAATGCAGCGGGGCCTCCTCCCCTGAATGTGAAGTAGAACGAGGA GACAGGGCAGAAGGGGCAGAAATAAAACAAGTGACAAAGCCAACAACCAGGGGCCTCAGCAGTAGCAGTGGC AGTGGTGGCAGCAGTAAGAGCCAAAGCAGCCTCCCCTGAGGGCCAGGCCTTGGAGAACCGGATGAAGCAGCTC TCCCTACAGTGCTCAAAGGGAAGAGATGGAATTATTGCTGACATAAAAATGGTGCAGATTGGC**TGA**TTCATCCTG GGCCCTGGCCGATGTGCATATCAACATTTATACATGGAACTGGAGAACATTGTGCCAATAATCATTTAATATATG CCAAATCTTACACGTCTACTCTAAACTGCTCTAATGAAGTTTCAGTGACCTTGAGGGCTAAAGATTGTTCTTCTG GGTAAGAGCTCTTGGGCTGGTTTTTCAGAGCAGAGTTCTTGTTGTGGGTAGACTGTGACTAGGTTCACAGCCTTT GTGGAACATTCCGTATAACGGCATTGTGGAAGCAATAACTAGTTCCTATGAAAGAACCAGAGCTGGGAAGATGGC CACCTCTGGTGTTCAGAGGGGGCTTTTAAAAAAGCGTGTATGCTGGGATACCCATTAAAACCATTTTCTAGAAGG CTACCATGAGCTGCACTTTTTGGGGTGGGAAAGGTGAATGCCAGTGGGGATGCGGGGGGATGAGGGTAGGAGGGA TATTCTAAGAGAAGTGCATGTGAAGAATGGTTGCCACTGTTATTAGATTGACAAGATGTTAATTTCTCTGTAGGT TGTAACTTTAAAATAAATGAAATTATTTAAGGGTTATGCTGCACTAGTATTCCTTAGAGGAAACAGTTCTTTAA AGTTAGGAAAGGGAGTAGGCAGGCATGTTTTGGCAAAGGCTGTTAATAGTAGTTAAGTGTTAAGACTGCTTTTCT TTAACGTTTTCATGGTAATGCATATTTAGAGCACTGTATTTTTGTCTTGTTTAAGAAAATTTAGCATTTCTAAAAG AAAAAAGCAACCCTCTTTCAAACTGTTAATTCTGTCACAGCCTGTATATTTTAGTCATTTGTAAATCTCTTCATA CAATAGTGACTTCTTTTTTGACTGATACAGTATCTTAATTACAAGGTTATTTTTGTACTTGTCTTAATACACTAAG TGTAATAAAAACGGCTTGAG

## 413/2825 FIGURE 377

MEQGLEEEEVDPRIQGELEKLNQSTDDINRRETELEDARQKFRSVLVEATVKLDELVKKIGKAVEDSKPYWEAR RVARQAQLEAQKATQDFQRATEVLRAAKETISLAEQRLLEDDKRQFDSAWQEMLNHATQRVMEAEQTKTRSELVH KETAARYNAAMGRMRQLEKKLKRAINKSKPYFELKAKYYVQLEQLKKTVDDLQAKLTLAKGEYKMALKNLEMISD EIHERRRSSAMGPRGCGVGAEGSSTSVEDLPGSKPEPDAISVASEAFEDDSCSNFVSEDDSETQSVSSFSSGPTS PSEMPDQFPAVVRPGSLDLPSPVSLSEFGMMFPVLGPRSECSGASSPECEVERGDRAEGAENKTSDKANNNRGLS SSSGSGGSSKSQSSTSPEGQALENRMKQLSLQCSKGRDGIIADIKMVQIG

### 414/2825 FIGURE 378

GACGCCATCCCTGTGACCATGACGATGTCCTCCATCGTGATGAAGGCAGAGACAGAGACATCCCGATCTGGGGA ACTCTGATCCAGTATATACGGCCTGTGTTCGTGTCCCGGTCAGACCAGGATTCTCGCAGGAAAACAGTAGAAGAA ATCAAGAGACGGGCGCAGTCCAACGGAAAGTGGCCACAGATAATGATTTTTCCAGAAGGAACTTGTACAAACAGG ACCTGCCTAATTACCTTCAAACCTGGTGCATTCATCCCTGGAGCGCCCGTCCAGCCTGTGGTTTTACGATATCCA AATAAACTGGACACCATCACATGGACGTGGCAAGGACCTGGAGCGCTGGAAATCCTGTGGCTCACGCTGTCAG TTTCACAACCAAGTGGAAATCGAGTTCCTTCCTGTGTACAGCCCTTCTGAGGAGGAGAAGAGGAACCCCGCGCTG  ${\tt TATGCCAGCAACGTGCGGCGAGTC} \underline{{\tt ATG}} {\tt GCCGAGGCCTTGGGTGTCTCCGTGACTACACGTTCGAGGACTGC}$ CAGCTGGCCCTGGCGGAAGGACAGCTCCGTCTCCCCGCTGACACTTGCCTTTTAGAATTTGCCAGGCTCGTGCGG GGCCTCGGGCTAAAACCAGAAAAGCTTGAAAAAGATCTGGACAGATACTCAGAAAGAGCCAGGATGAAGGGAGGA GAGAAGATAGGTATTGCGGAGTTTGCCGCCTCCCTGGAAGTCCCCGTTTCTGACTTGCTGGAAGACATGTTTTCA GCCCGGACCCTGGACACCATCCAGCTGGCTTTCAAGATGTACGGAGCGCAAGAGGACGGCAGCGTCGGCGAAGGT CAAGAGGAGAAGGGGAAGATCACATTCGCTGACTTCCACAGGTTTGCAGAAATGTACCCTGCCTTCGCAGAGGAA TACCTGTACCCGGATCAGACACATTTCGAAAGCTGTGCAGAGACCTCACCTGCGCCAATCCCAAACGGCTTCTGT GCCGATTTCAGCCCGGAAAACTCAGACGCTGGGCGGAAGCCTGTTCGCAAGAAGCTGGAT**TAG**GACCCAGGGTTG  ${\tt TCACTCCTGCTGTACAGGCACTGTCTTCAGCCCGAGTTCCAGGGGGCCTCGGGGGGCTGTTTGTATCTTGTTCCTTT}$ GTGAAGTGTGTGCAGAACCGACGCTTACTGTGCGAGAATCGGAGGGCGCGCACGCGGATCCCCCGCCTGGCCTG GACCCCGTGGGGTCAGGTTCCCTGCCGGGGGGGGGGCACCGGTGCCGCCCCGTGTTCTCCCACGGGGCCCTGGTT TCGAGTCTCTGTCACAGCCTCTTCCGGCGGCAGCGTGCACCGGGGGGGCCTCCGTGCACACTCAGCACACGCCTG CCACACAGCGTGCGCTTGCGTGTCACTCTGGCACGAAACCTGTCTGCCTCTGTGGATCCACAGCCTGGCAGAGCC GAGCCGTCACCTGATTTTTCAGTGTTTCTACCTGTGTGCTGGAGCTCATGAGTATTTTATAAACTCCATTTAGGT ACTTCAGGAAACATGCAGCATTTTTTAAAAAATGAAAATTGTTTTTCTACTTCATTTTTCCTTTTAGAGTCAAAG GATATTTATTATAGGCCTTTTTTTTTTTAATATAGAATCTGAGGCTGTTTGGGCTTTGACTTAAATTTCCATCA GGCCTCTCTCCAGCAGGTAATCCCTCTCCTTCCGCTGGGTCCCCTGGGGAGGTGTGAACTCAAGGGCCTAGCCCC AAAACACTTTTTCTGCTTTTCTTAATCCTTTTCCAGTCCCCTCTTTTTTTATAAACGTTGGCAGTTTGATGTTTC TGTTTCGGCATAACGTAATCCATTTCACTGTAGCCTAAACTCCAGTCCGAGGTTGGATATTGTTCAAATGAGCAG GGCCCGAGCTGGAAGCCCAAGGCAGCCGCCGCCGTGCCGCTCCTCCCTTGCCCTCAGGCCAGGTCCCTGCTGGAA GCGGCTGCATCTTCCTGTCAGCCCTGGTTTCCATGGTGACTGGCGTCACGCAGCCACCCGAGTATGGCTGACCTT ATGTCTGGTGACAGCCGTGCGGACACCACTCCTCTGCAGCACTGCCTCCCAGCGCCAGGGTCGCGGGCACATC CCACTGAGAGCGGGGTCCTGCCCCATCTTAGAGTCAAAGGCAGAGGGGCTTCCAGGCCCTGGATGGGGTATTTT CGAAGTCAATTTCAGGGCTCTTGGCTGTTGGACACAGAAGAAATCCTAGTGCAGCCTTTGGTAGCTAACAGTCAC TGATTTTATAATTGGAGAATGCGTAAAGATTCATTTTTCAAGGAGAAGAGCCTGCAAATGGCCAATGAAGGAGGT AAATAAACTAAGATATTCCGAGGGAAGGGACCCAGGCCACCTCCCTTCCGCAGGTCTGCAGATGAAGGGTTTTTT TGATTCTCTTTTCTCTTCGTGGCGACTTAGAGTTGGCGGATATTCGGAACTGTGAATGTACATAGCGTTGAGTTA  $\verb|AACCCCTTGTGTGAGACAGGACGCAGGCCCCTGGTGGCCTGGGGGGCCAGACCCGTGGGCAGGTGGGGCAT| \\$ AAAAAAAAAAAAAAAAAAAAA

## 415/2825 FIGURE 379

 $\label{thm:constraint} $$ MAEALGVSVTDYTFEDCQLALAEGQLRLPADTCLLEFARLVRGLGLKPEKLEKDLDRYSERARMKGGEKIGIAEF $$ AASLEVPVSDLLEDMFSLFDESGSGEVDLRECVVALSVVCRPARTLDTIQLAFKMYGAQEDGSVGEGDLSCILKT $$ ALGVAELTVTDLFRAIDQEEKGKITFADFHRFAEMYPAFAEEYLYPDQTHFESCAETSPAPIPNGFCADFSPENS DAGRKPVRKKLD$ 

### 416/2825 FIGURE 380

CCCCGTCTCCTGCCAGGCGCGGAGCCCTGCGAGCCGCGGGTGGGCCCCAGGCGCGCAGACATCGGCTCCGCC AAAGCGCGCTGGGCCGGGGCGCTGGGCGTCGCGGGGCTACTGTGCCGCTGTGCTGGGCGCTGTCATGATCGTG ATGGTGCCGTCGCTCATCAAGCAGCAGGTCCTTAAGAACGTGCGCATCGACCCCAGTAGCCTGTCCTTCAACATG TGGAAGGAGATCCCTATCCCCTTCTATCTCTCCGTCTACTTCTTTGACGTCATGAACCCCAGCGAGATCCTGAAG GGCGAGAAGCCGCAGGTGCGGGAGCCCCTACGTGTACAGGGAGTTCAGGCACAAAAGCAACATCACCTTC AACAACAACGACACCGTGTCCTTCCTCGAGTACCGCACCTTCCAGTTCCAGCCCTCCAAGTCCCACGGCTCGGAG AGCGACTACATCGTCATGCCCAACATCCTGGTCTTGGGTGCGGCGGTGATGATGGAGAATAAGCCCATGACCCTG AAGCTCATCATGACCTTGGCATTCACCACCCTCGGCGAACGTGCCTTCATGAACCGCACTGTGGGTGAGATCATG TGGGGCTACAAGGACCCCCTTGTGAATCTCATCAACAAGTACTTTCCAGGCATGTTCCCCCTTCAAGGACAAGTTC GGATTATTTGCTGAGCTCAACAACTCCGACTCTGGGCTCTTCACGGTGTTCACGGGGGTCCAGAACATCAGCAGG ATCCACCTCGTGGACAAGTGGAACGGGCTGAGCAAGGTTGACTTCTGGCATCCGATCAGTGCAACATGATCAAT GGAACTTCTGGGCAAATGTGGCCGCCCTTCATGACTCCTGAGTCCTCGCTGGAGTTCTACAGCCCGGAGGCCTGC CGATCCATGAAGCTAATGTACAAGGAGTCAGGGGTGTTTGAAGGCATCCCCACCTATCGCTTCGTGGCTCCCAAA ACCCTGTTTGCCAACGGGTCCATCTACCCACCCAACGAAGGCTTCTGCCCGTGCCTGGAGTCTGGAATTCAGAAC GTCAGCACCTGCAGGTTCAGTGCCCCCTTGTTTCTCTCCCATCCTCACTTCCTCAACGCCGACCCGGTTCTGGCA GAAGCGGTGACTGCCCTCACCCTAACCAGGAGGCACACTCCTTGTTCCTGGACATCCACCCGGTCACGGGAATC CCCATGAACTGCTCTGTGAAACTGCAGCTGAGCCTCTACATGAAATCTGTCGCAGGCATTGGACAAACTGGGAAG ATTGAGCCTGTGGTCCTGCTGTTTGCAGAGAGCGGGGCCATGGAGGGGGAGACTCTTCACACATTC TACACTCAGCTGGTGTTGATGCCCAAGGTGATGCACTATGCCCAGTACGTCCTCCTGGCGCTGGGCTGCGTCCTG CTGCTGGTCCTGTCATCTGCCAAATCCGGAGCCAAGTAGGTGCTGGCCAGAGGGCAGCCCGGGCTGACAGCCAT TCGCTTGCCTGCGGGGAAAGGGGCCTCAGATCGGACCCTCTGGCCAACCGCAGCCTGGAGCCCACCTCCAGCA CGTGAGCCAGCCAGGCCTGGCCGTGGCCTGACCGGCCCCCAGCCCCTACACCCCGCTTCTCCCGGACTCTCC CTGGCACACATACACATGCGTGCAGGCTTGTGCAGACACTCAGGGATGGAGCTGCTGCTGAAGGGACTTGTAG GGAGAGGCTCGTCAACAAGCACTGTTCTGGAACCTTCTCTCCACGTGGCCCACAGGCCTGACCACAGGGGCTGTG  ${\tt GGTCCTGCGTCCCCTTCCTCGGGTGAGCCTGGCCTGTCCCGTTCAGCCGTTGGGCCCAGGCTTCCTCCCCTCCAA}$ GGTGAAACACTGCAGTCCCGGTGTGGTGGCTCCCCATGCAGGACGGCCCAGGCTGGGAGTGCCGCCTTCCTGTGC CAAATTCAGTGGGGACTCAGTGCCCAGGCCCTGGCCACGAGCTTTGGCCTTGGTCTACCTGCCAGGCCAGGCAAA GCGCCTTTACACAGGCCTCGGAAAACAATGGAGTGAGCACAAGATGCCCTGTGCAGCTGCCCGAGGGTCTCCGCC CACCCGGCCGGACTTTGATCCCCCGGAGTCTTCACAGGCACTGCATCGGGTTGTCTGGCGCCCTTTTCCTCCA TGCCCCCACCCCTCACAGGGTCCCTCAGATTATAGGTGCCCAGGCTGAGGTGAAGAGGCCTGGGGGGCCCTGCCT TCCGGGCGCTCCTGGACCCTGGGGCAAACCTGTGACCCTTTTCTACTGGAATAGAAATGAGTTTTATCATCTTTG 

# FIGURE 381

MGCSAKARWAAGALGVAGLLCAVLGAVMIVMVPSLIKQQVLKNVRIDPSSLSFNMWKEIPIPFYLSVYFFDVMNP SEILKGEKPQVRERGPYVYREFRHKSNITFNNNDTVSFLEYRTFQFQPSKSHGSESDYIVMPNILVLGAAVMMEN KPMTLKLIMTLAFTTLGERAFMNRTVGEIMWGYKDPLVNLINKYFPGMFPFKDKFGLFAELNNSDSGLFTVFTGV QNISRIHLVDKWNGLSKVDFWHSDQCNMINGTSGQMWPPFMTPESSLEFYSPEACRSMKLMYKESGVFEGIPTYR FVAPKTLFANGSIYPPNEGFCPCLESGIQNVSTCRFSAPLFLSHPHFLNADPVLAEAVTGLHPNQEAHSLFLDIH PVTGIPMNCSVKLQLSLYMKSVAGIGQTGKIEPVVLPLLWFAESGAMEGETLHTFYTQLVLMPKVMHYAQYVLLA LGCVLLLVPVICQIRSQVGAGQRAARADSHSLACWGKGASDRTLWPTAAWSPPPAAVLRLCRSGSGHCWGLRSTL ASFACRVATTLPVLEGLGPSLGGGTGS

#### 418/2825 FIGURE 382

AGTCAAG**ATG**GAGGAGTACGCGCGAGAGCCTTGCCCATGGCGAATTGTGGATGACTGTGGTGGGGCCTTTACGAT GGGTACCATTGGTGGTGGTATCTTTCAAGCAATCAAAGGTTTTCGCAATTCTCCAGTGGGAGTAAACCACAGACT ACGAGGGAGTTTGACAGCTATTAAAACCAGGGCTCCACAGTTAGGAGGTAGCTTTGCAGTTTGGGGAGGGCTGTT TTCCATGATTGACTGTAGTATGGTTCAAGTCAGAGGAAAGGAAGATCCCTGGAACTCCATCACAAGTGGTGCCTT AACGGGAGCCATACTGGCAGCAAGAAATGGACCAGTGGCCATGGTTGGGTCAGCCGCAATGGGTGGCATTCTCCT AGCTTTAATTGAAGGAGCTGGTATCTTGTTGACAAGATTTGCCTCTGCACAGTTTCCCAATGGTCCTCAGTTTGC AGAAGACCCCTCCCAGTTGCCTTCAACTCAGTTACCTTCCTCACCTTTTGGAGACTATCGACAATATCAG**TAG**GA CTTCTTTCCTAGGATTTCTTTAACAGAACGAGTTGTGGTTCGAGAAGGATTTCAGAAGATCAAGTTACAGTCTGT TTTTAAAACCATAGGTGGGACAGCTATGGCCAATAGGCTATAAAGAGACATTTAGCACTTTTTTCTATTTAAAGG TGTTTAAATCGCTAAAGGAAAATACAGTAAGTGCTTGAAAGATGAAGGACCAAAAAGGCCCAAAAAACAGTGAAATA TGATCATCTCTTGCGGACTTCTCTGCCTGGTTTTGTGTGTTCTGTTATTCAAACAATAAAAAGCTGGTGGAA CTTACTCTTTCATAAGATAAGTTGTAGACTTCGATGTTTCATGCTCATGTACTTCAAATAATGCATGTTTTAT AGTTAGTCCCTCATCACTTGAAGTGACTTCTGAGAATTATGCAGAGTCAACATGGATCATTTCACAGTGAGATGC TTTATGGATTGAAGGATATGGTAAAATGTTTATAGTTTACTTTGAAAGTAAAATATACTATGTCTTGGTTTTGAG GATATTGGATACAAAACTCTCTTTCCTTTAGGGCTACTGAGTCTTGATTCCTGATCATCAGAAATTTCACCAGAAA CAACTTGCTTCCAATATACCCAATTCTATATGAAGAATTCATGGAGAGTGTACTGGCACTGGAAGAGTTTAGTGT TTCTTGTATGCTTGAAAATAAAGTATGTACTGTTTTGAATGTGTTCCAAGTCCTCTGCATAAACGATGTATTTTG GGGTCTGGTTGGGCCTGGAAAATGGATGAGCACTTCAGAACAGGTCATTTTCCTGATATTGGAAGTGACATGTGG  $\verb|CCCTATAGGAGGCATGATGTTAGTTAATTACACATTTGCCTACATCTGTGGGAAATGGAGAACAAAGCCATGTGG|$ 

## 419/2825 FIGURE 383

 ${\tt MEEYAREPCPWRIVDDCGGAFTMGTIGGGIFQAIKGFRNSPVGVNHRLRGSLTAIKTRAPQLGGSFAVWGGLFSMIDCSMVQVRGKEDPWNSITSGALTGAILAARNGPVAMVGSAAMGGILLALIEGAGILLTRFASAQFPNGPQFAEDPSQLPSTQLPSSPFGDYRQYQ$ 

#### 420/2825 FIGURE 384

**ATG**GCGACGGAGCAGAGGCCTTTCCACCTGGTGTTTCGGCGCGCTCTGGCTTCACCGGCCAGTTCGTGACCGAG GAGGTGGCCCGGGAGCAGCCGGAGCGGACCCAGCCTGCCCTGGGCGTGGCGGGCCGCTCCCGGGAGAAG  $\tt CTGCAGCGGGTGCTGGAGAAGGCGGCCCTGAAGCTGGGAAGACCAACACTGTCATCTGAAGTTGGAATCATCATC$ TGTGATATTGCTAATCCAGCCTCGCTTGATGAAATGGCTAAACAGGCAACAGTTGTCCTCAATTGCGTAGGACCA TATCGGTTTTATGGAGAACCTGTAATAAAAGCATGTATTGAAAATGGAGCCAGTTGTATCGACATCAGTGGAGAA  $\verb|CCTCAGTTTCTGGAACTAATGCAACTGAAGTATCATGAGAAAGCTGCAGACAAAGGGGTTTATATCATTGGAAGC| \\$ GTGGAAAGTTTCCTGACTATACATTCAGGACCTGAGGGGTTGAGCATTCATGATGGTACCTGGAAGTCAGCAATT TATGGTTTTGGAGATCAGAGTAATTTGAGAAAACTAAGAAATGTATCAAATCTGAAACCTGTCCCGCTCATTGGT CCAAAATTGAAGAGAAGGTGGCCAATTTCTTATTGTCGGGAACTCAAAGGTTATTCCATTCCTTTTATGGGATCT GATGTGTCTGTTGTAAGGAGGACTCAACGTTACTTGTATGAAAATTTAGAGGAATCACCAGTTCAGTATGCTGCG TTTGGAATTGGAAGGCAACTTCTCATAAAATTCCCATGGTTCTTCTCCTTTTGGCTATTTTTTCAAAACAAGGCCCA ACACAAAAACAGATTGATGCTGCCTCATTCACGCTGACATTCTTTGGTCAAGGATACAGCCAAGGCACTGGTACA TCTGAAGTC**TAA**ACACTGGAAGAATTAACTGAAGTCATAACGTGCGTGAATTAACAGCTTCTCTATTTGATATTT AGGAAATTGTCCTAGCTTACCCTAAATTTCAAATCTGAGTTGATTTTGTGATTTTATTGCTTATAACAGAGAACT CATATTTGACATATTTTTTCATTGATGTGTTCCTGGTAGATTTTCACGAATGAGCTGGCAGGTCTAATGGGGGA GGCGGCGTCCCAGTCTGTTGCAGCAGCATTCTCATCGGGGGTGCGCACACCATCGTTACTGTCGGGCAGTAAC AACTGCACTTTTTGCTGTTTTTGCCTAGTTTTTCGCCAATCTACACTGATTTTGGACTGTTACCTAAGTTGAAAA ATAAAAGGTTGTCAATCGAATGGTGGTTAATGTTGGACCTGCCGATGTATTTGTATAGTGGTAGAAACATGCTGC TTAAGTGGCCTAACCTGTTTCTTGCCAATAAGTAGGCTTATCATTTTATTTTTACGTAATTCTATATCTGTGACT AGGTTTTTAAGGATACAGCTTATAAGTTGCTATCAATTTTCACTACCTAAGCAGAATTTTTCTCTAATTTACTTT TTGTATTTTAACTAGGTTTTACATGGAAGCCCTAAAATAAGGCAAAAGACTTTTTCTTTTGTAATAAGCATATAA TAAACACGTATATACATAGCAAATTGA

## 421/2825 FIGURE 385

MATEQRPFHLVVFGASGFTGQFVTEEVAREQVDPERTQPALGVAGRSREKLQRVLEKAALKLGRPTLSSEVGIII
CDIANPASLDEMAKQATVVLNCVGPYRFYGEPVIKACIENGASCIDISGEPQFLELMQLKYHEKAADKGVYIIGS
SGFDSIPADLGVIYTRNKMNGTLTAVESFLTIHSGPEGLSIHDGTWKSAIYGFGDQSNLRKLRNVSNLKPVPLIG
PKLKRRWPISYCRELKGYSIPFMGSDVSVVRRTQRYLYENLEESPVQYAAYVTVGGITSVIKLMFAGLFFLFFVR
FGIGRQLLIKFPWFFSFGYFSKQGPTQKQIDAASFTLTFFGQGYSQGTGTDKNKPNIKICTQVKGPEAGYVATPI
AMVQAAMTLLSDASHLPKAGGVFTPGAAFSKTKLIDRLNKHGIEFSVISSSEV

## 422/2825 FIGURE 386

## 423/2825 FIGURE 387

MTERRVPFSLLRGPSWDPFRDWYPHSRLFDQAFGLPRLPEEWSQWLGGSSWPGYVRPLPPAAIESPAVAAPAYSR ALSRQLSSGVSEIRHTADRWRVSLDVNHFAPDELTVKTKDGVVEITGKHEERQDEHGYISRCFTRKYTLPPGVDP TQVSSSLSPEGTLTVEAPMPKLATQSNEITIPVTFESRAQLGGPEAAKSDETAAK

#### 424/2825 FIGURE 388

ACTGCGACTCGAGACAGCGCCCGGCAGGACAGCTCCAGA ATG AAAATGCGGTTCTTGGGGTTGGTGTTTTGAGTGAAATTATCTCTTACTGGGGATTCCCTAGTGAGGAATACCTAGTTGAGACAGAAGATGGATATATTCTGTG CCTTAACCGAATTCCTCATGGGAGGAAGAACCATTCTGACAAAGGTCCCAAACCAGTTGTCTTCCTGCAACATGG CTTGCTGGCAGATTCTAGTAACTGGGTCACAAACCTTGCCAACAGCAGCCTGGGCTTCATTCTTGCTGATGCTGG TTTTGACGTGTGGATGGGCAACAGCAGAGGAAATACCTGGTCTCGGAAACATAAGACACTCTCAGTTTCTCAGGA TGAATTCTGGGCTTTCAGTTATGATGAGATGGCAAAATATGACCTACCAGCTTCCATTAACTTCATTCTGAATAA AACTGGCCAAGAACAAGTGTATTATGTGGGTCATTCTCAAGGCACCACTATAGGTTTTATAGCATTTTCACAGAT CCCTGAGCTGGCTAAAAGGATTAAAATGTTTTTTGCCCTGGGTCCTGTGGCTTCCGTCGCCTTCTGTACTAGCCC TATGGCCAAATTAGGACGATTACCAGATCATCTCATTAAGGACTTATTTGGAGACAAAGAATTTCTTCCCCAGAG TGCGTTTTTGAAGTGGCTGGGTACCCACGTTTGCACTCATGTCATACTGAAGGAGCTCTGTGGAAATCTCTGTTT TCTTCTGTGTGGATTTAATGAGAGAAATTTAAATATGTCTAGAGTGGATGTATATACAACACATTCTCCTGCTGG AACTTCTGTGCAAAACATGTTACACTGGAGCCAGGCTGTTAAATTCCAAAAGTTTCAAGCCTTTGACTGGGGAAG CAGTGCCAAGAATTATTTTCATTACAACCAGAGTTATCCTCCCACATACAATGTGAAGGACATGCTTGTGCCGAC TGCAGTCTGGAGCGGGGTCACGACTGGCTTGCAGATGTCTACGACGTCAATATCTTACTGACTCAGATCACCAA CTTGGTGTTCCATGAGAGCATTCCGGAATGGGAGCATCTTGACTTCATTTGGGGCCTGGATGCCCCTTGGAGGCT TATATCCAAGAAAATGATAACTTTGAAGATGCCCAGTTCACTCTAGTTTCAATTAGAAACATACTAGCTATTTTT TCTTTAATTAGGGCTGGAATAGGAAGCCAGTGTCTCAACCATAGTATTGTCTCTTTAAGTCTTTTAAATATCACT GATGTGTAAAAAGGTCATTATATCCATTCTGTTTTTAAAATTTAAAATATTGACTTTTTGCCCTTCATAGGAC AAAGTAATATATGTGTTGGAATTTTAAAATTGTGTTGTCATTGGTAAATCTGTCACTGACTTAAGCGAGGTATAA AAGTACGCAGTTTTCATGTCCTTGCCTTAAAGAGCTCTCTAGTCTAACGGTCTTGTAGTTAGAGATCTAAATGAC ATTTTATCATGTTTTCCTGCAGCAGGTGCATAGTCAAATCCAGAAATATCACAGCTGTGCCAGTAATAAGGATGC TAACAATTAATTTTATCAAACCTAACTGTGACAGCTGTGATTTGACACGTTTTAATTGCTCAGGTTAAATGAAAT AGTTTTCCGGCGTCTTCAAAAACAAATTGCACTGATAAAACAAAAACAAAAGTATGTTTTAAATGCTTTGAAGAC TGATACACTCAACCATCTATATTCATGAGCTCTCAATTTCATGGCAGGCCATAGTTCTACTTATCTGAGAAGCAA ATCCCTGTGGAGACTATACCACTATTTTTTCTGAGATTAATGTACTCTTGGAGCCCGCTACTGTCGTTATTGATC ACATCTGTGTGAAGCCAAAGCCCCGTGGTTGCCCATGAGAAGTGTCCTTGTTCATTTTCACCCAAATGAAGTGTG AACGTGATGTTTTCGGATGCAAACTCAGCTCAGGGATTCATTTTGTGTCTTAGTTTTATATGCATCCTTATTTTT AATACACCTGCTTCACGTCCCTATGTTGGGAAGTCCATATTTGTCTGCTTTTCTTGCAGCATCATTTCCTTACAA TACTGTCCGGTGGACAAAATGACAATTGATATGTTTTTCTGATATAATTACTTTAGCTGCACTAACAGTACAATG CTTGTTAATGGTTAATATAGGCAGGGCGAATACTACTTTGTAACTTTTAAAGTCTTAAACTTTTCAATAAAATTG AGTGAGACTTATAGGCCC

## 425/2825 FIGURE 389

MKMRFLGLVVCLVLWPLHSEGSGGKLTAVDPETNMNVSEIISYWGFPSEEYLVETEDGYILCLNRIPHGRKNHSD KGPKPVVFLQHGLLADSSNWVTNLANSSLGFILADAGFDVWMGNSRGNTWSRKHKTLSVSQDEFWAFSYDEMAKY DLPASINFILNKTGQEQVYYVGHSQGTTIGFIAFSQIPELAKRIKMFFALGPVASVAFCTSPMAKLGRLPDHLIK DLFGDKEFLPQSAFLKWLGTHVCTHVILKELCGNLCFLLCGFNERNLNMSRVDVYTTHSPAGTSVQNMLHWSQAV KFQKFQAFDWGSSAKNYFHYNQSYPPTYNVKDMLVPTAVWSGGHDWLADVYDVNILLTQITNLVFHESIPEWEHL DFIWGLDAPWRLYNKIINLMRKYQ

#### 426/2825 FIGURE 390

CGCAGGCTGGAAGGAAGACGAACCTACGAAGCAGAGATCTGAAGACAGC**ATG**TACACAGCCATTCCCCAGAGTGG CTCTCCATTCCCAGGCTCAGTGCAGGATCCAGGCCTGCATGTGTGGCGGGTGGAGAAGCTGAAGCCGGTGCCTGT CAACACGCTGCTGGGAGAGCGGCCTGTGCAGCACCGCGAGGTGCAGGGCAATGAGTCTGACCTCTTCATGAGCTA CTTCCCACGGGGCCTCAAGTACCAGGAAGGTGGTGTGGAGTCAGCATTTCACAAGACCTCCACAGGAGCCCCAGC TGCCATCAAGAAACTCTACCAGGTGAAGGGGAAGAAGAACATCCGTGCCACCGAGCGGGCACTGAACTGGGACAG CTTCAACACTGGGGACTGCTTCATCCTGGACCTGGGCCAGAACATCTTCGCCTGGTGTGGTGGAAAGTCCAACAT CCTGGAACGCAACAAGGCGAGGGACCTGGCCCTGGCCATCCGGGACAGTGAGCGACAGGGCCAAGGCCCAGGTGGA GATTGTCACTGATGGGGAGGAGCCTGCTGAGATGATCCAGGTCCTGGGCCCCAAGCCTGCTCTGAAGGAGGGCAA CCCTGAGGAAGACCTCACAGCTGACAAGGCAAATGCCCAGGCCGCAGCTCTGTATAAGGTCTCTGATGCCACTGG ACAGATGAACCTGACCAAGGTGGCTGACTCCAGCCCCTTTGCCCTTGAACTGCTGATATCTGATGACTGCTTTGT CCTGCAGGTGGCCGAGGGCTTCATCTCGCGCATGCAGTACGCCCCGAACACTCAGGTGGAGATTCTGCCTCAGGG  ${\tt CCGTGAGAGTCCCATCTTCAAGCAATTTTTCAAGGACTGGAAA}{{\tt TGA}}{\tt GGGTGGGCGTCTTCCTGCCCCATGCTCCC}$  $\tt CTGCCCCCACCACCTGCTTGCTTCTCTGGCTGCCTGGTCAGTGCAGAGGTGCCCCCTGCAGATGTTCAAT$ AAAGGAGACAAGTGCTTTCCC

## 427/2825 FIGURE 391

MYTAIPQSGSPFPGSVQDPGLHVWRVEKLKPVPVAQENQGVFFSGDSYLVLHNGPEEVSHLHLWIGQQSSRDEQG ACAVLAVHLNTLLGERPVQHREVQGNESDLFMSYFPRGLKYQEGGVESAFHKTSTGAPAAIKKLYQVKGKKNIRA TERALNWDSFNTGDCFILDLGQNIFAWCGGKSNILERNKARDLALAIRDSERQGKAQVEIVTDGEEPAEMIQVLG PKPALKEGNPEEDLTADKANAQAAALYKVSDATGQMNLTKVADSSPFALELLISDDCFVLDNGLCGKIYIWKGRK ANEKERQAALQVAEGFISRMQYAPNTQVEILPQGRESPIFKQFFKDWK

# FIGURE 392

CCGGTCCTGAGACACGCTGTGTGGCTGAAAAGTGAAGGCAAGAGCTCATTTGGCCTCTGTGCTCCCCTCCGCAAG GGATCGTTTCTCCAGAAGAGCTGGATATTCTTTCGCCCAGTT**ATC**GCAGACAAGTTAACGAGAATTGCTATTGTC AACCATGACAAATGTAAACCTAAGAAATGTCGACAGGAATGCAAAAAGAGTTGTCCTGTAGTTCGAATGGGAAAA TTATGCATAGAGGTTACACCCCAGAGCAAAATAGCATGGATTTCCGAAACTCTTTGTATTGGTTGTGGTATCTGT ATTAAGAAATGCCCCTTTGGCGCCTTATCAATTGTCAATCTACCAAGCAACTTGGAAAAAGAAACCACACATCGA TATTGTGCCAATGCCTTCAAACTTCACAGGTTGCCTATCCCTCGTCCAGGTGAAGTTTTGGGATTAGTTGGAACT AATGGTATTGGAAAGTCAGCTGCTTTAAAAATTTTAGCAGGAAAACAAAAGCCAAACCTTGGAAAGTACGATGAT CCTCCTGACTGGCAGGAGATTTTGACTTATTTCCGTGGATCTGAATTACAAAATTACTTTACAAAGATTCTAGAA GATGACCTAAAAGCCATCATCAAACCTCAATATGTAGCCAGATTCCTAAGGCTGGCAAAGGGGACAGTGGGATCT ATTTTGGACCGAAAAGATGAAACAAAGACACAGGCAATTGTATGTCAGCAGCTTGATTTAACCCACCTAAAAGAA CGAAATGTTGAAGATCTTTCAGGAGGAGAGTTGCAGAGATTTGCTTGTGCTGTCGTTTGCATACAGAAAGCTGAT ATTTTCATGTTTGATGAGCCTTCTAGTTACCTAGATGTCAAGCAGCGTTTAAAGGCTGCTATTACTATACGATCT CTAATAAATCCAGATAGATATATCATTGTGGTGGAACATGATCTAAGTGTATTAGACTATCTCTCCGACTTCATC TGCTGTTTATATGGTGTACCAAGCGCCTATGGAGTTGTCACTATGCCTTTTAGTGTAAGAGAAGGCATAAACATT TTTTTGGATGGCTATGTTCCAACAGAAAACTTGAGATTCAGAGATGCATCACTTGTTTTTAAAGTGGCTGAGACA GCAAATGAAGAAGTTAAAAAGATGTGTATGTATAAATATCCAGGAATGAAGAAAAAAATGGGAGAATTTGAG CTAGCAATTGTAGCTGGAGAGTTTACAGATTCTGAAATTATGGTGATGCTGGGGGAAAATGGAACGGGTAAAACG ACATTTATCAGAATGCTTGCTGGAAGACTTAAACCTGATGAAGGAGGAGAAGTACCAGTTCTAAATGTCAGTTAT AAGCCACAGAAAATTAGTCCCAAATCAACTGGAAGTGTTCGCCAGTTACTACATGAAAAGATAAGAGATGCTTAT ACTCACCCACATTTGTGACCGATGTAATGAAGCCTCTGCAAATTGAAAACATCATTGATCAAGAGGTGCAGACA TTATCTGGTGGTGAACTACAGCGAGTACGTTTACGCCTTTGCTTGGGCAAACCTGCTGATGTCTATTTAATTGAT GAACCATCTGCATATTTGGATTCTGAGCAAAGACTGATGGCAGCTCGAGTTGTCAAACGTTTCATACTCCATGCA AAAAAGACAGCCTTTGTTGTGGAACATGACTTCATCATGGCCACCTATCTAGCGGATCGCGTCATCGTTTTTGAT GAACAAAAGAAGAGTGGAAACTACTTTTTCTTGGATGAT**TAG**ACTGACTCTGAGAATATTGATAAGCCATTTATT AAAAGGAGTATTTACTAGAATTTTTTGTCATATAAAACTTGAATCAGGATTTTATGCCCCACATACTCTGGAACT TGAAGTATAATATATATAATAAAAAAGCCAGTTGGGTTCTAAATTGTAGTTGAAACACAGAAAATGCCA CTTTTCTGTTCCTGAAGAGGCTCTTTTGTGCATAATATTCTAAAATGAAGACATTTCAAGCTATACAAATTACTT CCAAGTTTTCATGATGTATGGGAAGATTTTCAGTAGGTGTATTATATTCACGGTACCAAATGCTGACCAGTGTTG CTCCATTTTTTAAATCTTGAAAAGGGTTTCTGTACTTACCTGGTTTGCCAAGTATGCCAGTGTAATGAAACTGCC GTTTTCAGTCTTTGCACTATATTACCAGTATATGGTTTCCGAGGAAGATTATCTACTGCAAAACACCACTGTTGG AAAAATAGGTATTTTTAAATTGTTTTTAATCCTTTTTTGGTGCTTTTTAAACATGTTTAAGCAAAAACCAATTCAG GTCATTCTCTAGGTAATGTCATTTTTGTACACATATATTTATATAATCACTGATTGAGATTTAGGAAAAAGCATT TCTAAAGAATATTTGCTTCCCTTAGAACTACAGACTCGAAATCTTTAAAGATGGTGCCTAAGCATCTATGTATTT TTTTTAAGTTCCACAGATTTTTCTGTTGGGCAGGCCAAGGATTATAAACCACTTCCCTAAAGGCAACATTAATGC AAAAGTCCCCAGATGGCAATACAAAGTATCCCCTGGTACCACATATATTCATTTGTGAGTTTTGGATATAGAGCAC ATTATCTAAACCATTTTGTAGTTCCAAAAACCCATCTAAATTTCTTGAGTTCCTGAATTTTGAACAGGATTACCT ATGACTGGGGCAAGCAAAATTGAGGAGGAAATTAGAAACTGTTTGACAAACTTTAAGAGCTACTTGAAATAACAG AAGTCTTGATTAATATGCCAAATAATGGCTAGAAAGTATGGTTTAACTGGACCCTATTATGCCTTTTAAAAATAAT TTCAGTAACCCATAAATACATGTTGTAAAAAATTCAAATATACAGAATGGAATAAAAAAATGATCTCCCTTTATT ACCCTCCCAAAGGTTACCAGCGTTTGAATTTAATAATGTATATTCTTTCATGCTTTTTTCTGTGCACTTACCTAA GTGTGAATATGTAAAGGGTTTGTTTTGTATACAAATGGGATTATACTAAAATAAGTAATGCCTATTTTTAAGGAT AGGTTAAATTTGTGAATGATCATTTCAAATATATTGAATAAAATAAGCAAAAGCTATTGTTATTTACTGATCCTG 

#### 429/2825 FIGURE 393

MADKLTRIAIVNHDKCKPKKCRQECKKSCPVVRMGKLCIEVTPQSKIAWISETLCIGCGICIKKCPFGALSIVNL PSNLEKETTHRYCANAFKLHRLPIPRPGEVLGLVGTNGIGKSAALKILAGKQKPNLGKYDDPPDWQEILTYFRGS ELQNYFTKILEDDLKAIIKPQYVARFLRLAKGTVGSILDRKDETKTQAIVCQQLDLTHLKERNVEDLSGGELQRF ACAVVCIQKADIFMFDEPSSYLDVKQRLKAAITIRSLINPDRYIIVVEHDLSVLDYLSDFICCLYGVPSAYGVVT MPFSVREGINIFLDGYVPTENLRFRDASLVFKVAETANEEEVKKMCMYKYPGMKKKMGEFELAIVAGEFTDSEIM VMLGENGTGKTTFIRMLAGRLKPDEGGEVPVLNVSYKPQKISPKSTGSVRQLLHEKIRDAYTHPQFVTDVMKPLQ IENIIDQEVQTLSGGELQRVRLRLCLGKPADVYLIDEPSAYLDSEQRLMAARVVKRFILHAKKTAFVVEHDFIMA TYLADRVIVFDGVPSKNTVANSPQTLLAGMNKFLSQLEITFRRDPNNYRPRINKLNSIKDVEQKKSGNYFFLDD

#### 430/2825 **FIGURE 394**

# 431/2825 FIGURE 395

 ${\tt MSHKQIYYSDKYDDEEFEYRHVMLPKDIAKLVPKTHLMSESEWRNLGVQQSQGWVHYMIHEPEPHILLFRRPLPKKPKK}$ 

#### 432/2825 FIGURE 396

 $\tt AGCCAGAAATGTGAAGTGCTAGCTGAAGGATGAGCAGCTAGCCAGGCAAAGGGGGGCA{\color{red} ATC} {\color{blue} GCGGCTTCCTGT}$ GTTCTACTGCACACTGGGCAGAAGATGCCTCTGATTGGTCTGGGTACCTGGAAGAGTGAGCCTGGTCAGGTAAAA ATTGGGGAGGCCCTGAAGGACGTGGGACCAGGCAAGGCGGTGCCTCGGGAGGAGCTGTTTGTGACATCCAAG CTGTGGAACACCAAGCACCCCGAGGATGTGGAGCCTGCCCTCCGGAAGACTCTGGCTGACCTCCAGCTGGAG TATCTGGACCTGTACCTGATGCACTGGCCTTATGCCTTTGAGCGGGGAGACACCCCTTCCCCAAGAATGCTGAT GGGACTATATGCTACGACTCCACCCACTACAAGGAGACTTGGAAGGCTCTGGAGGCACTGGTGGCTAAGGGGCTG GTGCAGGCGCTGGGCCTGTCCAACTTCAACAGTCGGCAGATTGATGACATACTCAGTGTGGCCTCCGTGCGTCCA GCTGTCTTGCAGGTGGAATGCCACCCATACTTGGCTCAAAATGAGCTAATTGCCCACTGCCAAGCACGTGGCTTG GAGGTAACTGCTTATAGCCCTTTGGGCTCCTCTGATCGTGCATGGCGTGATCCTGATGAGCCTGTCCTGCTGGAG GAACCAGTAGTCCTGGCATTGGCTGAAAAGTATGGCCGATCTCCAGCTCAGATCTTGCTCAGGTGGCAGGTCCAG CGGAAAGTGATCTGCATCCCCAAAAGTATCACTCCTTCTCGAATCCTTCAGAACATCAAGGTGTTTGACTTCACC TTTAGCCCAGAAGAGATGAAGCAGCTAAATGCCCTGAACAAAAATTGGAGATATATTGTGCCTATGCTTACGGTG  ${\tt GATGGGAAGAGTCCCAAGGGATGCAGGGCATCCTCTGTACCCCTTTAATGACCCGTAC{\color{red}{\bf TGA}}{\tt GACCACAGCTTC}}$ TTGGCCTCCCTTCCAGCTCTGCAGCTAATGAGGTCCTGCCACAACGGAAAGAGGGAGTTAATAAAGCCATTGGAG CATCCAT

## 433/2825 FIGURE 397

MAASCVLLHTGQKMPLIGLGTWKSEPGQVKAAVKYALSVGYRHIDCAAIYGNEPEIGEALKEDVGPGKAVPREEL FVTSKLWNTKHHPEDVEPALRKTLADLQLEYLDLYLMHWPYAFERGDNPFPKNADGTICYDSTHYKETWKALEAL VAKGLVQALGLSNFNSRQIDDILSVASVRPAVLQVECHPYLAQNELIAHCQARGLEVTAYSPLGSSDRAWRDPDE PVLLEEPVVLALAEKYGRSPAQILLRWQVQRKVICIPKSITPSRILQNIKVFDFTFSPEEMKQLNALNKNWRYIV PMLTVDGKRVPRDAGHPLYPFNDPY

## 434/2825 FIGURE 398

## 435/2825 FIGURE 399

MEALPLLAATTPDHGRHRRLLLLPLLLFLLPAGAVQGWETEERPRTREEECHFYAGGQVYPGEASRVSVADHSLH LSKAKISKPAPYWEGTAVIDGEFKELKLTDYRGKYLVFFFYPLDFTFVCPTEIIAFGDRLEEFRSINTEVVACSV DSQFTHLAWINTPRRQGGLGPIRIPLLSDLTHQISKDYGVYLEDSGHTLRGLFIIDDKGILRQITLNDLPVGRSV DETLRLVQAFQYTDKHGEVCPAGWKPGSETIIPDPAGKLKYFDKLN

## 436/2825 FIGURE 400

CCGCTGGGCGTAGCTGCGACTCGGCGGAGTCCCGGCGGCGCGTCCTTGTTCTAACCCGGCGCGCCATGACCGTCG CGCGGCCGAGCGTGCCCGCGCGCGCTCCTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGCTGTTGTGCC GTTTTCCCGAGGATACTGTAATAACGTACAAATGTGAAGAAGCTTTGTGAAAATTCCTGGCGAGAAGGACTCAG TGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGCTGCGAGGTGCCAACAAGGC TAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTATTTTCCAGTCGGTACTGTTGTGGAATATGAGT CAGTCGAATTTTGTAAAAAGAAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTG GCATATTATTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGTTTTT GTCTTATTTCAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAAAATTTATTGTCCAGCACCAC CACAAATTGACAATGGAATAATTCAAGGGGAACGTGACCATTATGGATATAGACAGTCTGTAACGTATGCATGTA ATAAAGGATTCACCATGATTGGAGAGCACTCTATTTATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCC CACCACCTGAATGCAGAGGAAAATCTCTAACTTCCAAGGTCCCACCAACAGTTCAGAAACCTACCACAGTAAATG GGAGTACACCTGTTTCCAGGACAACCAAGCATTTTCATGAAACAACCCCAAATAAAGGAAGTGGAACCACTTCAG GTACTACCCGTCTTCTATCTGGGCACACGTGTTTCACGTTGACAGGTTTGCTTGGGACGCTAGTAACCATGGGCT TGCTGACT**TAG**CCAAAGAAGAGTTAAGAAGAAAATACACACAAGTATACAGACTGTTCCTAGTTTCTTAGACTTA TCTGCATATTGGATAAAATAAATGCAATTGTGCTCTTCATTTAGGATGCTTTCATTGTCTTTAAGATGTGTTAGG AATGTCAACAGAGCAAGGAGAAAAAAGGCAGTCCTGGAATCACATTCTTAGCACACCTACACCTCTTGAAAATAG AACAACTTGCAGAATTGAGAGTGATTCCTTTCCTAAAAGTGTAAGAAAGCATAGAGATTTGTTCGTATTTAGAAT GGGATCACGAGGAAAAGAGAAGTGATTTTTTTCCACAAGATCTGTAATGTTATTTCCACTTATAAAGGAA ATAAAAAATGAAAAACATTATTTGGATATCAAAAGCAAATAAAAACCCAATTCAGTCTCTTCTAAGCAAAATTGC TAAAGAGAGATGAACCACATTATAAAGTAATCTTTGGCTGTAAGGCATTTTCATCTTTCCTTCGGGTTGGCAAAA TATTTTAAAGGTAAAACATGCTGGTGAACCAGGGGTGTTGATGGTGATAAGGGGAGGAATATAGAATGAAAGACTG AATCTTCCTTTGTTGCACAAATAGAGTTTGGAAAAAGCCTGTGAAAGGTGTCTTCTTTGACTTAATGTCTTTAAA AGTATCCAGAGATACTACAATATTAACATAAGAAAAGATTATATTATTTCTGAATCGAGATGTCCATAGTCAA GT

## 437/2825 FIGURE 401

MTVARPSVPAALPLLGELPRLLLLVLLCLPAVWGDCGLPPDVPNAQPALEGRTSFPEDTVITYKCEESFVKIPGE KDSVICLKGSQWSDIEEFCNRSCEVPTRLNSASLKQPYITQNYFPVGTVVEYECRPGYRREPSLSPKLTCLQNLK WSTAVEFCKKKSCPNPGEIRNGQIDVPGGILFGATISFSCNTGYKLFGSTSSFCLISGSSVQWSDPLPECREIYC PAPPQIDNGIIQGERDHYGYRQSVTYACNKGFTMIGEHSIYCTVNNDEGEWSGPPPECRGKSLTSKVPPTVQKPT TVNVPTTEVSPTSQKTTTKTTTPNAQATRSTPVSRTTKHFHETTPNKGSGTTSGTTRLLSGHTCFTLTGLLGTLV TMGLLT

#### 438/2825 FIGURE 402

**ATG**TTCCGAGCGGCGGCTCCGGGGGCGGCCTCATTGCTACGATTTCAGAGTACCCTGGTAATA GCTGAGCATGCAAATGATTCCCTAGCACCCCATTACTTTAAATACCATTACTGCAGCCACACGCCTTGGAGGTGAA GTGTCCTGCTTAGTAGCTGGAACCAAATGTGACAAGGTGGCACAAGATCTCTGTAAAGTAGCAGGCATAGCAAAA  ${\tt GTTCTGGTGGCTCAGCATGATGTTGCAAAAGGCCTACTTCCAGAGGAACTGACACCATTGATTTTGGCAACTCAG}$ AAGCAGTTCAATTACACACATCTGTGCTGGAGCATCTGCCTTCGGAAAGAACCTTTTGCCCAGAGTAGCAGCC AAACTTGAGGTTGCCCCGATTTCTGACATCATTGCAATCAAGTCACCTGACACATTTGTGAGAACTATTTATGCA  ${\tt GGAAATGCTCTATGTACAGTGAAGTGTGATGAGAAAGTGAAAGTGTTTTCTGTCCGTGGAACATCCTTTGATGCT}$ GCAGCAACAAGTGGCGGTAGTGCCAGTTCAGAAAAGGCATCAAGTACTTCACCAGTGGAAATATCAGAGTGGCTT GACCAGAAATTAACAAAAAGTGATCGACCAGAGCTAACAGGTGCCAAAGTGGTGGTATCTGGTGGTCGAGGCTTG AAGAGTGGAGAGAACTTTAAGTTGTTATATGACTTGGCAGATCAACTACATGCTGCAGTTGGTGCTTCCCGTGCT GCTGTTGATGCTGGCTTTGTTCCCAATGACATGCAAGTTGGACAGACGGGAAAAATAGTAGCACCAGAACTTTAT ATTGCTGTTGGAATATCTGGAGCCATCCAACATTTAGCTGGGATGAAAGACAGCAAGACAATTGTGGCAATTAAT  ${\tt ATGACTGAGATATTGAAGAAAAA} {\tt {\tt TGA}} {\tt ATCAGGATCATGCCTTAAAAAGAAACTTTTGTTAAAGTATTCCACTG}$ AAATCACAGATATTTGTGGGTATTATAACAATCATTGGAAAGCATGGAGAGCTACATTTCATAATTTGAGGGAAA ATTTCTAACAGATGCCAGAATGCTTGTTTATGGGATTGCTGTTTTCCTTTTAATTATTTGTGGTTCCAAACAAT TATTGTTTGAACTTTTTTAATTCTGTACTAAAATCTATAATAAAGCTTTTCCACAGCTTTAAAACT

## 439/2825 FIGURE 403

MFRAAAPGQLRRAASLLRFQSTLVIAEHANDSLAPITLNTITAATRLGGEVSCLVAGTKCDKVAQDLCKVAGIAK VLVAQHDVYKGLLPEELTPLILATQKQFNYTHICAGASAFGKNLLPRVAAKLEVAPISDIIAIKSPDTFVRTIYA GNALCTVKCDEKVKVFSVRGTSFDAAATSGGSASSEKASSTSPVEISEWLDQKLTKSDRPELTGAKVVVSGGRGL KSGENFKLLYDLADQLHAAVGASRAAVDAGFVPNDMQVGQTGKIVAPELYIAVGISGAIQHLAGMKDSKTIVAIN KDPEAPIFQVADYGIVADLFKVVPEMTEILKKK

#### 440/2825 FIGURE 404

**ATG**GAGCTGTGCGGGCTGCCCCGGCCGCCCATGCTGCTGCTGCTGTTGGCGACACTGCTGCCGCCG ATGTTGGCGCTGCTGACTCAGGTGGCGCTGGTGGTGCAGGTGGCGGAGGCGCCCCGAGCGTCTCGGCC AAGCCGGGGCCGCTGTGGCCCCTGCCGCTCTCGGTGAAGATGACCCCGAACCTGCTGCATCTCGCCCCGGAG AACTTCTACATCAGCCACAGCCCCAATTCCACGGCGGGCCCCTCCTGCACCCTGCTGGAGGAAGCGTTTCGACGA TATCATGGCTATATTTTTGGTTTCTACAAGTGGCATCATGAACCTGCTGAATTCCAGGCTAAAACCCAGGTTCAG CAACTTCTTGTCTCAATCACCCTTCAGTCAGAGTGTGATGCTTTCCCCAACATATCTTCAGATGAGTCTTATACT TTACTTGTGAAAGAACCAGTGGCTGTCCTTAAGGCCAACAGAGTTTGGGGAGCATTACGAGGTTTAGAGACCTTT AGCCAGTTAGTTATCAAGATTCTTATGGAACTTTCACCATCAATGAATCCACCATTATTGATTCTCCAAGGTTT TCTCACAGAGGAATTTTGATTGATACATCCAGACATTATCTGCCAGTTAAGATTATTCTTAAAACTCTGGATGCC ATGGCTTTTAATAAGTTTAATGTTCTTCACTGGCACATAGTTGATGACCAGTCTTTCCCATATCAGAGCATCACT TTTCCTGAGTTAAGCAATAAAGGAAGCTATTCTTTGTCTCATGTTTATACACCAAATGATGTCCGTATGGTGATT GAATATGCCAGATTACGAGGAATTCGAGTCCTGCCAGAATTTGATACCCCTGGGCATACACTATCTTGGGGAAAA GGTCAGAAAGACCTCCTGACTCCATGTTACAGTAGACAAAACAAGTTGGACTCTTTTGGACCTATAAACCCTACT TTGGGAGGAGATGAAGTGGAATTTAAATGTTGGGAATCAAATCCAAAAATTCAAGATTTCATGAGGCAAAAAGGC TTTGGCACAGATTTTAAGAAACTAGAATCTTTCTACATTCAAAAGGTTTTTGGATATTATTGCAACCATAAACAAG GGATCCATTGTCTGGCAGGAGGTTTTTGATGATAAAGCAAAGCTTGCGCCGGGCACAATAGTTGAAGTATGGAAA GACAGCGCATATCCTGAGGAACTCAGTAGAGTCACAGCATCTGGCTTCCCTGTAATCCTTTCTGCTCCTTGGTAC TTAGATTTGATTAGCTATGGACAAGATTGGAGGAAATACTATAAAGTGGAACCTCTTGATTTTGGCGGTACTCAG AAACAGAAACATTTTCATTGGTGGAGAAGCTTGTCTATGGGGAGAATATGTGGATGCAACTAACCTCACTCCA AGATTATGGCCTCGGGCAAGTGCTGTTGGTGAGAGACTCTGGAGTTCCAAAGATGTCAGAGATATGGATGACGCC TATGACAGACTGACAAGGCACCGCTGCAGGATGGTCGAACGTGGAATAGCTGCACAACCTCTTTATGCTGGATAT TGTAACCATGAGAACATG**TAA**AAAATGGAGGGGAAAAAGGCCACAGCAATCTGTACTACAATCAACTTTATTTTG 

## 441/2825 FIGURE 405

MELCGLGLPRPPMLLALLLATLLAAMLALLTQVALVVQVAEAARAPSVSAKPGPALWPLPLSVKMTPNLLHLAPE
NFYISHSPNSTAGPSCTLLEEAFRRYHGYIFGFYKWHHEPAEFQAKTQVQQLLVSITLQSECDAFPNISSDESYT
LLVKEPVAVLKANRVWGALRGLETFSQLVYQDSYGTFTINESTIIDSPRFSHRGILIDTSRHYLPVKIILKTLDA
MAFNKFNVLHWHIVDDQSFPYQSITFPELSNKGSYSLSHVYTPNDVRMVIEYARLRGIRVLPEFDTPGHTLSWGK
GQKDLLTPCYSRQNKLDSFGPINPTLNTTYSFLTTFFKEISEVFPDQFIHLGGDEVEFKCWESNPKIQDFMRQKG
FGTDFKKLESFYIQKVLDIIATINKGSIVWQEVFDDKAKLAPGTIVEVWKDSAYPEELSRVTASGFPVILSAPWY
LDLISYGQDWRKYYKVEPLDFGGTQKQKQLFIGGEACLWGEYVDATNLTPRLWPRASAVGERLWSSKDVRDMDDA
YDRLTRHRCRMVERGIAAQPLYAGYCNHENM

#### 442/2825 FIGURE 406

 $\tt CTGGTCCCGAGCAGGTGTGAGGGGATTCACTTGTGTGCGGAACTCCTCGGAACC \textbf{ATG} \tt GCGTCCCTTTCCCTT$ GCCATCGCCATTGGAGACTTGGTAAAGAGCACCTTGGGACCCAAAGGCATGGACAAAATTCTTCTAAGCAGTGGA CGAGATGCCTCTCTTATGGTAACCAATGATGGTGCCACTATTCTAAAAAACATTGGTGTTTGACAATCCAGCAGCT AAAGTTTTAGTTGATATGTCAAGGGTTCAAGATGATGAAGTTGGTGATGGCACTACCTCTGTTACCGTTTTAGCA GCAGAATTATTAAGGGAAGCAGAATCTTTAATTGCAAAAAAGATTCATCCACAGACCATCATAGCGGGTTGGAGA GAAGCCACGAAGGCTGCAAGAGAGGCGCTGTTGAGTTCTGCAGTTGATCATGGTTCCGATGAAGTTAAATTCCGT CAAGATTTAATGAATATTGCGGGCACAACATTATCCTCAAAACTTCTTACTCATCACAAAGACCACTTTACAAAG TTAGCTGTAGAAGCAGTTCTCAGACTGAAAGGCTCTGGCAACCTGGAGGCAATTCATATTATCAAGAAGCTAGGA GGAAGTTTGGCAGATTCCTATTTAGATGAAGGCTTCCTGTTGGATAAAAAAATTGGAGTAAATCAACCAAAACGA ATTGAAAATGCTAAAATTCTTATTGCAAATACTGGTATGGATACAGACAAAATAAAGATATTTGGTTCCCGGGTA AGAGTTGACTCTACAGCAAAGGTTGCAGAAATAGAACATGCGGAAAAAGGAAAAATGAAGGAGAAAGTTGAACGT GCTGGTGTCATGGCTATTGAGCATGCAGATTTTGCAGGTGTGGAACGCCTAGCTCTTGTCACAGGTGGTGAAATT GCCTCTACCTTTGATCACCCAGAACTGGTGAAGCTTGGAAGTTGCAAACTTATCGAGGAAGTCATGATTGGAGAA GACAAACTCATTCACTTTTCTGGGGTTGCCCTTGGTGAGGCTTGTACCATTGTTTTGCGTGGTGCCACTCAACAA ATTTTAGATGAAGCAGAAAGATCATTGCATGATGCTCTTTGTGTTCTTGCGCAAACTGTAAAGGACTCTAGAACA GTTTATGGAGGAGGCTGTTCTGAGATGTTGATGGCTCATGCTGTGACACAGCTTGCCAATAGAACACCAGGCAAA GAAGCTGTTGCAATGGAGTCTTATGCTAAAGCACTGAGAATGTTGCCAACCATCATAGCTGACAATGCAGGCTAT GACAGTGCAGACCTGGTGGCACAGCTCAGGGCTGCTCACAGTGAAGGCAATACCACTGCTGGATTGGATATGAGG GAAGGCACCATTGGAGATATGGCTATCCTGGGTATAACAGAAAGTTTTCAAGTGAAGCGACAGGTTCTTCTGAGT GCAGCTGAAGCAGCAGAGGTGATTCTGCGTGTGGACAACATCATCAAAGCGGCACCCAGGAAACGTGTCCCTGAT CACCACCCTGT**TAA**GCATTCCCACGTGCTGTCGATCTTTGGACCAGTTTCTAGCAAAGTTGTGTTTGAAAGATA CTCTATTAAAGAAGACTGTGGAATCTGTTTATCGGTGCCCATTATATCCTTAAGTTTTGGATATTTAGCTGACCTT CGCTTTAACATAGGTCTAATTTATTTGCCGTGTCATTTTCCATACAAATCAGTTGATTTAAAGGAGTTCATTTCG 

## 443/2825 FIGURE 407

MASLSLAPVNIFKAGADEERAETARLTSFIGAIAIGDLVKSTLGPKGMDKILLSSGRDASLMVTNDGATILKNIG VDNPAAKVLVDMSRVQDDEVGDGTTSVTVLAAELLREAESLIAKKIHPQTIIAGWREATKAAREALLSSAVDHGS DEVKFRQDLMNIAGTTLSSKLLTHHKDHFTKLAVEAVLRLKGSGNLEAIHIIKKLGGSLADSYLDEGFLLDKKIG VNQPKRIENAKILIANTGMDTDKIKIFGSRVRVDSTAKVAEIEHAEKEKMKEKVERILKHGINCFINRQLIYNYP EQLFGAAGVMAIEHADFAGVERLALVTGGEIASTFDHPELVKLGSCKLIEEVMIGEDKLIHFSGVALGEACTIVL RGATQQILDEAERSLHDALCVLAQTVKDSRTVYGGGCSEMLMAHAVTQLANRTPGKEAVAMESYAKALRMLPTII ADNAGYDSADLVAQLRAAHSEGNTTAGLDMREGTIGDMAILGITESFQVKRQVLLSAAEAAEVILRVDNIIKAAP RKRVPDHHPC

#### 444/2825 FIGURE 408A

GAATCCAAGGGGGCCAGTTCCTGCCGTCTTCTGCCTCTTGATCTCCGCCACCGTCTTCAGGCCAGGCCTT GGATGGTATACTGTAAATTCAGCATATGGAGATACCATTATCATACCTTGCCGACTTGACGTACCTCAGAATCTC AGTGTGCAGTACGACGATGTACCAGAATACAAAGACAGATTGAACCTCTCAGAAAACTACACTTTGTCTATCAGT AATGCAAGGATCAGTGATGAAAAGAGATTTGTGTGCATGCTAGTAACTGAGGACAACGTGTTTGAGGCACCTACA ATAGTCAAGGTGTTCAAGCAACCATCTAAACCTGAAATTGTAAGCAAAGCACTGTTTCTCGAAACAGAGCAGCTA AAAAAGTTGGGTGACTGCATTTCAGAAGACAGTTATCCAGATGGCAATATCACATGGTACAGGAATGGAAAAGTG CTACATCCCCTTGAAGGAGCGGTGGTCATAATTTTTAAAAAGGAAATGGACCCAGTGACTCAGCTCTATACCATG ACTTCCACCCTGGAGTACAAGACAACCAAGGCTGACATACAAATGCCATTCACCTGCTCGGTGACATATTATGGA CCATCTGGCCAGAAAACAATTCATTCTGAACAGGCAGTATTTGATATTTACTATCCTACAGAGCAGGTGACAATA CAAGTGCTGCCACCAAAAAATGCCATCAAAGAAGGGGATAACATCACTCTTAAATGCTTAGGGAATGGCAACCCT CCCCCAGAGGAATTTTTGTTTTACCTACCAGGACAGCCCGAAGGAATAAGAAGCTCAAATACTTACACACTGACG GATGTGAGGCGCAATGCAACAGGAGACTACAAGTGTTCCCTGATAGACAAAAAAAGCATGATTGCTTCAACAGCC ATCACAGTTCACTATTTGGATTTGTCCTTAAACCCAAGTGGAGAAGTGACTAGACAGATTGGTGATGCCCTACCC GTGTCATGCACAATATCTGCTAGCAGGAATGCAACTGTGGTATGGATGAAAGATAACATCAGGCTTCGATCTAGC CCGTCATTTTCTAGTCTTCATTATCAGGATGCTGGAAACTATGTCTGCGAAACTGCTCTGCAGGAGGTTGAAGGA CTAAAGAAAAGAGTCATTGACTCTCATTGTAGAAGGCAAACCTCAAATAAAAATGACAAAGAAAACTGATCCC AGTGGAAGCGTCATAAACCAAACAGAGGAATCTCCTTATATTAATGGCAGGTATTATAGTAAAATTATCATTTCC CCTGAAGAGAATGTTACATTAACTTGCACAGCAGAAAACCAACTGGAGAGAACAGTAAACTCCTTGAATGTCTCT GCTATAAGTATTCCAGAACACGATGAGGCAGACGAGATAAGTGATGAAAAACAGAGAAAAAGGTGAATGACCAGGCA AAGAAGTCAAAGACTGCATCAAAACATGTAAACAAGGACCTCGGTAATATGGAAGAAACAAAAAGTTAGAAGAA AACAATCACAAAACTGAAGCCTAAGAGAAAACTGTCCTAGTTGTCCAGAGATAAAAATCATATAGACCAATTGA AGCATGAACGTGGATTGTATTTAAGACATAAACAAAGACATTGACAGCAATTCATGTTCAAGTATTAAGCAGTTC GAGGATTTCTTTTGGTTTGCCTTTTATGTAAATTTTTTACGTAGCTATTTTTATACACTGTAAGCTTTGTTCTG TTTCTGATTCTGATTGCTATCAGCAATGCCCCAAACTTTCTCATAAGCACCTAAAACCCAAAGGTGGCAGCTTGT TAGACACATACATAATGGTACTCCCAAACTGACAATTTTACCTATTCTGAAAAAGACATAAAACAGAATTTGG TAGCACTTACCTCTACAGACACCTGCTAATAAATTATTTTCTGTCAAAAGAAAAAACACAAGCATGTGTGAGAGA CAGTTTGGAAAAATCATGGTCAACATTCCCATTTTCATAGATCACAATGTAAATCACTATAATTACAAATTGGTG TTTTGTTAACTACCCTACAGATATTGAATGCACCTTGAGATAATTTAGTGTTTTAACTGATACATAATTTATCAA GCAGTACATGAAAGTGTAATAAAAATGTCTATGTATCTTTAGTTACATTCAAATTTGTAACTTTATAAACATG TTTTATGCTTGAGGAAATTTTTAAGGTGGTAGTATAAATGGAAACTTTTTGAAGTAGACCAGATATGGGCTACTT GTGACTAGACTTTTAAACTTTGCTCTTTCAAGCAGAAGCCTGGTTTCTGGGAGAACACTGCACAGTGATTTCTTT CCCAGGATTTACACAACTTTAAAGGGAAGATAAATGAACATCAGATTTCTAGGTATAGAACTATGTTATTGAAAG ATTAAAGTCAGATGAGAATAGGAATAATACTTTGCCACTTCTGCATTATTTAGAAACATACGTTATTGTACATTT GTAAACCATTTACTGTCTGGGCAATAGTGACTCCGTTTAATAAAAGCTTCCGTAGTGCATTGGTATGGATTAAAT GCATAAAATATCTTAGACTCGATGCTGTATAAAATATTATGGGAAAAAAGAAATACGTTATTTTGCCTCTAAACT TTTATTGAAGTTTTATTTGGCAGGAAAAAAATTGAATCTTGGTCAACATTTAAACCAAAGTAAAAGGGGAAAAA 

## 445/2825 FIGURE 408B

#### 446/2825 FIGURE 409A

TCAACACAGGACAATGCAAGCCCATGAGCTGTTCCGGTATTTTCGAATGCCAGAGCTGGTTGACTTCCGACAGTA CGTGCGTACTCTTCCGACCAACACGCTTATGGGCTTCGGAGCTTTTGCAGCACTCACCACCTTCTGGTACGCCAC AAGATCCGCACTACTTGACAGCGACGAGCCCTTGGTGTATTTCTATGATGATGTCACAACATTATACGAAGGTTT CCAGAGGGGAATACAGGTGTCAAATAATGGCCCTTGTTTAGGCTCTCGGAAACCAGACCAACCCTATGAATGGCT TTCATATAAACAGGTTGCAGAATTGTCGGAGTGCATAGGCTCAGCACTGATCCAGAAGGGCTTCAAGACTGCCCC AGATCAGTTCATTGGCATCTTTGCTCAAAATAGACCTGAGTGGGTGATTATTGAACAAGGATGCTTTGCTTATTC GATGGTGATCGTTCCACTTTATGATACCCTTGGAAATGAAGCCATCACGTACATAGTCAACAAAGCTGAACTCTC TCTGGTTTTTGTTGACAAGCCAGAGAAGGCCAAACTCTTATTAGAGGGTGTAGAAAATAAGTTAATACCAGGCCT TAAAATCATAGTTGTCATGGATGCCTACGGCAGTGAACTGGTGGAACGAGGCCAGAGGTGTGGGGTGGAAGTCAC CAGCATGAAGGCGATGGAGGACCTGGGAAGAGCCAACAGACGGAAGCCCAAGCCTCCAGCACCTGAAGATCTTGC TCTCGCCCATATGTTTGAGAGAGTTGTAGAGTGTGTAATGCTGTGTCATGGAGCTAAAATCGGATTTTTCCAAGG AGATATCAGGCTGCTCATGGATGACCTCAAGGTGCTTCAACCCACTGTCTTCCCCGTGGTTCCAAGACTGCTGAA GAAAGAAGCAGAGCTTCGCAGCGGCATCATCAGAAACAACAGCCTGTGGGACCGGCTGATCTTCCACAAAGTACA GTCGAGCCTGGGCGGAAGAGTCCGGCTGATGGTGACAGGAGCCGCCCGGTGTCTGCCACTGTGCTGACGTTCCT GCCTGGAGACTGGACCGCAGGCCATGTTGGGGCCCCGATGCCGTGCAATTTGATAAAACTTGTTGATGTGGAAGA GGACCCAGCGAAAACAGCAGAAGCTTTGGACAAAGACGGCTGGTTACACACAGGGGACATTGGAAAATGGTTACC AAATGGCACCTTGAAAATTATCGACCGGAAAAAGCACATATTTAAGCTGGCACAAGGAGAATACATAGCCCCTGA AAAGATTGAAAATATCTACATGCGAAGTGAGCCTGTTGCTCAGGTGTTTGTCCACGGAGAAAGCCTGCAGGCATT TGAGGAACTGTGCAGAAATAAGGATGTCAAAAAAGCTATCCTCGAAGATATGGTGAGACTTGGGAAGGATTCTGG TCTGAAACCATTTGAACAGGTCAAAGGCATCACATTGCACCCTGAATTATTTTCTATCGACAATGGCCTTCTGAC CATAGGAATGCTAGAGGAAACAGAACACCGCCTTACAGTCACCTCATGTTGCAGACCATGTTTATGGTAATACAC ACTTTCCAAAATGAGCCTTAAAAATTGTAAAGGGGATACTATAAATGTGCTAAGTTATTTGAGACTTCCTCAGTT TAAAAAGTGGGTTTTAAATCTTCTGTCTCCCTGCTTTTCTAATCAAGGGGTTAGGACTTTGCTATCTCTGAGATG TCTGCTACTTGCTGCAAATTCTGCAGCTGTCTGCTGCTCTAAAGAGTACAGTGCACTAGAGGGAAGTGTTCCCTT TAAAAATAAGAACAACTGTCCTGGCTGGAGAATCTCACAAGCGGACCAGAGATCTTTTTAAATCCCTGCTACTGT CCCTTCTCACAGGCATTCACAGAACCCTTCTGATTCGTAAGGGTTACGAAACTCATGTTCTTCTCCAGTCCCCTG TGGTTTCTGTTGGAGCATAAGGTTTCCAGTAAGCGGGAGGGCAGATCCAACTCAGAACCATGCAGATAAGGAGCC TCTGGCAAATGGGTGCTCATCAGAACGCGTGGATTCTCTTTCATGGCAGAATGCTCTTGGACTCGGTTCTCCAGG CCTGATTCCCCGACTCCATCCTTTTTCAGGGGTTATTTAAAAATCTGCCTTAGATTCTATAGTGAAGACAAGCAT TTCAAGAAAGAGTTACCTGGATCAGCCATGCTCAGCTGTGACGCCTGAATAACTGTCTACTTTATCTTCACTGAA CCACTCACTCTGTGTAAAGGCCAACAGATTTTTAATGTGGTTTTCATATCAAAAGATCATGTTGGGATTAACTTG CCTTTTTCCCCAAAAAATAAACTCTCAGGCAAGCATTTCTTTAAAGCTATTAAGGGAGTATATACTTGAGTACTT ATTGAAATGGACAGTAATAAGCAAATGTTCTTATAATGCTACCTGATTTCTATGAAATGTGTTTGACAAGCCAAA ATTCTAGGATGTAGAAATCTGGAAAGTTCATTTCCTGGGATTCACTTCTCCAGGGATTTTTTAAAGTTAATTTGG GAAATTAACAGCAGTTCACTTTATTGTGAGTCTTTTGCCACATTTGACTGAATTGAGCTGTCATTTGTACATTTAA AGCAGCTGTTTTGGGGTCTGTGAGAGTACATGTATTATATACAAGCACAACAGGGCTTGCACTAAAGAATTGTCA TTGTAATAACACTACTTGGTAGCCTAACTTCATATATGTATTCTTAATTGCACAAAAAGTCAATAATTTGTCACC TTGGGGTTTTGAATGTTTGCTTTAAGTGTTTGGCTATTTCTATGTTTTATAAACCAAAACAAAATTTCCAAAAACA

# 447/2825 **FIGURE 409B**

ATGAAGGAAACCAAAATAAATATTTCTGCATTTCG

#### 448/2825 FIGURE 410

MQAHELFRYFRMPELVDFRQYVRTLPTNTLMGFGAFAALTTFWYATRPKPLKPPCDLSMQSVEVAGSGGARRSAL LDSDEPLVYFYDDVTTLYEGFQRGIQVSNNGPCLGSRKPDQPYEWLSYKQVAELSECIGSALIQKGFKTAPDQFI GIFAQNRPEWVIIEQGCFAYSMVIVPLYDTLGNEAITYIVNKAELSLVFVDKPEKAKLLLEGVENKLIPGLKIIV VMDAYGSELVERGQRCGVEVTSMKAMEDLGRANRRKPKPPAPEDLAVICFTSGTTGNPKGAMVTHRNIVSDCSAF VKATENTVNPCPDDTLISFLPLAHMFERVVECVMLCHGAKIGFFQGDIRLLMDDLKVLQPTVFPVVPRLLNRMFD RIFGQANTTLKRWLLDFASKRKEAELRSGIIRNNSLWDRLIFHKVQSSLGGRVRLMVTGAAPVSATVLTFLRAAL GCQFYEGYGQTECTAGCCLTMPGDWTAGHVGAPMPCNLIKLVDVEEMNYMAAEGEGEVCVKGPNVFQGYLKDPAK TAEALDKDGWLHTGDIGKWLPNGTLKIIDRKKHIFKLAQGEYIAPEKIENIYMRSEPVAQVFVHGESLQAFLIAI VVPDVETLCSWAQKRGFEGSFEELCRNKDVKKAILEDMVRLGKDSGLKPFEQVKGITLHPELFSIDNGLLTPTMK AKRPELRNYFRSQIDDLYSTIKV

#### 449/2825 FIGURE 411

GAGAATGGCTCAACTCTCCAAATAGAGAGAATTTCTATCAGTTGCAAGTACGAAAATTTCCTGCCGATTATATAA AATACTGGGAGTTTGCAGTTTATCTGGAAGAATGTGAACTGGCTAAACAGCTTTATCCAAAGGAAAACGATTTGG TGTTTTTAGCTCCTGAGAGAATAAATGAAGAGAAGAAGATACAGAGAATGACATACAAGATCTCCACGAAT TCCAGACTCAAGAGAACTTTCCGGCCAATTTAAACGAACTTGTGAATTGTAATCAGTTCTCTGGTAACTA CACAAAGGAAGTTGAAAGCCATGTCTCTGTTGGGTAGTCGGAACCAACTGGCTAGAGCTGTTCTGAATCCAAACC CTATGGACTTCTGTACAAAAGATTTACTGACTACAACATCTGAGAGAATTATTGCGTACTTAAGAGATTTCAATG AAGATCAAAAGAAAGCAATAGAAACTGCATATGCTATGGTGAAACACTCACCATCAGTTGCCAAAATCTGCTTGA TTCATGGACCACCTGGAACAGGAAAATCAAAAACTATTGTTGGCCTCCTCTATCGTCTACTGACAGAGAACCAGA CAGCTGTTGATGAACTCATGAAAAAAATTATCCTTGAATTCAAAGAAAAATGTAAAGACAAGAAGAATCCTTTAG GAAACTGTGGAGATATAAATTTAGTACGACTGGGTCCAGAAAAGTCTATTAATAGTGAGGTTCTAAAGTTCAGTT TTCTAGATTATCAGCTGGATGAGCTTTCCCGGCAGCGAGCTCTATGCCGAGGTGGACGGGAAATACAGAGGCAAG AATTAGATGAAAACATTTCCAAAGTTTCTAAGGAAAGGCAGGAACTTGCTTCTAAAATTAAAGAGGTTCAAGGAC GCCCACAGAAAACACAGAGTATCATCTTAGAGTCCCATATCATCTGCTGCACGTTGAGCACAAGTGGTGGTT TACTACTTGAGTCTGCTTTCCGTGGGCAAGGGGGTGTCCCCTTCAGCTGTCATTGTTGATGAGGCTGGACAGT CTTGTGAAATTGAGACTCTTACTCCACTCATCCATCGCTGCAATAAGCTCATCCTAGTAGGAGATCCTAAGCAGC TCCCTCCGACAGTCATCTCTATGAAAGCACAGGAGTATGGCTACGACCAGTCAATGATGGCTCGCTTCTGCAGAC TGCTGGAAGAGATGTAGAACACATGATCAGCAGGCTGCCCATTCTACAGCTCACTGTTCAGTACAGGATGC ATCCAGACATATGCCTCTTCCCTTCTAATTATGTTTATAACAGAAACTTAAAAAACAAATAGACAGAAGACCA TTCGATGTTCATCAGATTGGCCATTTCAGCCATACCTTGTGTTTGATGTTGGAGATGGTTCAGAAAGACGGGATA ATGACTCATATATAAATGTTCAAGAAATAAAACTGGTGATGGAAATAATTAAGCTTATTAAAGACAAAAGAAAAGG ATGTTAGTTTTCGAAACATTGGCATAATAACTCATTACAAGGCCCAGAAGACGATGATTCAGAAGGATTTGGACA AAGAGTTCGATAGAAAAGGACCAGCAGAAGTAGACACTGTGGATGCATTCCAGGGTCGGCAGAAGGATTGTGTTA TTGTTACGTGTGTCAGAGCAAATAGCATCCAAGGTTCAATTGGATTCCTGGCAAGTTTGCAGAGATTGAATGTCA CCATCACACGAGCCAAGTACAGCCTCTTCATCCTCGGACATTTGAGGACCCTGATGGAAAACCAGCATTGGAATC AGCTGATTCAGGATGCTCAGAAGCGTGGTGCCATTATTAAGACCTGTGACAAAAACTATAGACATGATGCAGTGA  $\verb|CCCAGGGTGGTTTGCCCAGCAGCAAGCTAGACAGTTGGATTTGCCAAGACATCTGTTGCTGCTTCTCTATACCACA| \\$ CACCCTCTGACTCCAAGGAAATTACTCTTACTGTTACTTCAAAGGACCCTGAAAGACCTCCTGTTCATGACCAAC TTCAGGACCCACGACTGCTGAAGAGGATGGGCATTGAGGTCAAAGGAGGAATATTCCTTTGGGATCCACAACCCT ATATACAGCAGCCCGCTGCTGTAGTGGCTGCTCTGAGCAGCCACAAACCTCCCGTGCGGGGGGAACCTCCAGCTG CCAGTCCCGAGGCTTCCACGTGTCAGAGCAAATGTGATGACCCGGAAGAGGAGCTCTGTCACAGGAGAGAGGCCA GGGCTTTCAGTGAAGGGGAGCAGGAGAAGTGTGGTTCCGAGACCCATCACACCAGGAGGAACTCTAGGTGGGACA GCAGCCACAGCATATTGTAAACTGAAGATGACCAGCTCGTGGGACCATCTAGATAAGCTTGTTTTTTGTAAGGAG  $\verb|TTTGTGTGCTGTTGGAAAACATGGAAAATGCATCCTTAACACCTGAGCCTCTGGTCATCTTCAGTATTTTCTGTC|\\$ ATTTGCAAAAGCTTTCAGAGGGCATTGTGTATCCGTAATAATGTCCTTGAAGTCAGAGACTGGAAATGTTGATCT CTTAGTCTTCTATAGACAAGGAGGTCAGACTGGAGTGAATGTTGTAAAAGGTTGAAATGTATATTTGTATAGCAAA TAACAAAATCCTTTTAAAATTT

## 450/2825 FIGURE 412

MRNGKTECYLSIQTQENFPANLNELVNCIVISSLVTTQRKLKAMSLLGSRNQLARAVLNPNPMDFCTKDLLTTTS
ERIIAYLRDFNEDQKKAIETAYAMVKHSPSVAKICLIHGPPGTGKSKTIVGLLYRLLTENQRKGHSDENSNAKIK
QNRVLVCAPSNAAVDELMKKIILEFKEKCKDKKNPLGNCGDINLVRLGPEKSINSEVLKFSLDSQVNHRMKKELP
SHVQAMHKRKEFLDYQLDELSRQRALCRGGREIQRQELDENISKVSKERQELASKIKEVQGRPQKTQSIIILESH
IICCTLSTSGGLLLESAFRGQGGVPFSCVIVDEAGQSCEIETLTPLIHRCNKLILVGDPKQLPPTVISMKAQEYG
YDQSMMARFCRLLEENVEHNMISRLPILQLTVQYRMHPDICLFPSNYVYNRNLKTNRQTEAIRCSSDWPFQPYLV
FDVGDGSERRDNDSYINVQEIKLVMEIIKLIKDKRKDVSFRNIGIITHYKAQKTMIQKDLDKEFDRKGPAEVDTV
DAFQGRQKDCVIVTCVRANSIQGSIGFLASLQRLNVTITRAKYSLFILGHLRTLMENQHWNQLIQDAQKRGAIIK
TCDKNYRHDAVKILKLKPVLQRSLTHPPTIAPEGSRPQGGLPSSKLDSGFAKTSVAASLYHTPSDSKEITLTVTS
KDPERPPVHDQLQDPRLLKRMGIEVKGGIFLWDPQPSSPQHPGATPPTGEPGFPVVHQDLSHIQQPAAVVAALSS
HKPPVRGEPPAASPEASTCQSKCDDPEEELCHRREARAFSEGEQEKCGSETHHTRRNSRWDKRTLEQEDSSSKKR
KLL

#### 451/2825 FIGURE 413A

GCGGCGCTGGGTCGGTGGCGAGGCTGAGGAGAAGGAGGAGCGGCCGTGGAGGCTTCGCCGCCTAGGTACTGCT  $\tt ATAACCAGAATTTGGTAGAAAAAGGATTTACTTGTTGGGGGCCCTCTTGATAAAAAGAG{\color{red} \underline{ATG}} \\ \texttt{TGGGGGGGATTCTCG} \\$ ACCTGCTAACAGAACTGGACCTTTTCGTGGGAGCCAAGAAGAAAGGTTTGCTCCCGGGTGGAACAGGGATTATCC TCCTCCTCCCCTTAAGAGTCATGCTCAAGAGAGACACTCTGGCAACTTTCCTGGCAGAGATTCACTTCCCTTTGA TTTCCAGGGGCATTCGGGGCCTCCTTTTGCAAATGTAGAGGAGCATTCTTTCAGCTATGGAGCTAGAGACGGACC ACCTATGGACTATAGGGGTGGAGATGGTACTTCTATGGATTATAGAGGTAGGGAGGCACCTCATATGAACTACAG AGACAGGGATGCTCACGCTGTTGACTTCAGAGGTAGGGATGCTCCTCCATCTGACTTCAGGGGCCGGGGCACTTA TGATTTAGATTTTAGAGGCCGGGATGGATCCCATGCAGATTTTAGGGGAAGGGATTTATCAGATTTTGGATTTTAG GGCCAGAGAACAGTCCCGTTCTGATTTTAGGAATAGAGATGTATCTGATTTGGACTTTAGAGACAAAGACGGAAC ACAAGTAGACTTTAGAGGCCGAGGTTCAGGTACTACTGATCTAGACTTTAGGGACAGGGATACGCCACATTCAGA TTTCAGAGGTAGACACCGATCTAGGACTGATCAGGATTTTAGGGGCAGAGAGATGGGATCTTGTATGGAATTTAA AGATAGGGAGATGCCCCTGTGGATCCAAATATTTTTGGATTACATTCAGCCCTCTACACAAGATAGAGAACATTC TGGTATGAATGTGAACAGGAGAGAAGAATCCACACATGACCATACGATAGAAAGGCCTGCTTTTGGCATTCAGAA GGGAGAATTTGAGCATTCAGAAACAAGAGAGAGAGAAACACAAGGTGTAGCCTTTGAACATGAGTCTCCAGCAGA CTTTCAGAACAGCCAAAGTCCAGTTCAAGACCAAGATAAGTCACAGCTTTCTGGACGTGAAGAGCAGAGTTCAGA TGCTGGTCTGTTTAAAGAAGAAGCCGTCTGGACTTTCTTGGGCGGCAAGACACCGATTACAGAAGCATGGAGTA CCGTGATGTGGATCATAGGCTGCCAGGAAGCCAGATGTTTGGCTATGGCCAGAGCAAGTCTTTTCCAGAGGGCAA AACTGCCCGAGATGCCCAACGGGACCTTCAGGATCAAGATTATAGGACCGGCCCAAGTGAGGAGAAACCCAGCAG GCTTATTCGATTAAGTGGGGTACCTGAAGATGCCACAAAAGAAGAGATTCTTAATGCTTTTCGGACTCCTGATGG CATGCCTGTAAAGAACTTGCAGTTGAAGGAGTATAACACAGGTTACGACTATGGCTATGTCTGCGTGGAGTTTTC ACTCTTGGAAGATGCCATCGGATGCATGGAGGCCAACCAGGGAACTCTAATGATCCAGGACAAAGAAGTTACCCT GGAGTATGTATCAAGCCTGGATTTTTGGTACTGCAAACGATGTAAGGCAAACATTGGTGGGCACCGATCTTCCTG ATCCATACCAGCACCATTGGAAAAACAGCCCAACCAGCCCCTAAGACCAGCTGATAAGGAACCTGAACCCAGGAA AACTATCATGCTAAAGCGTATCTATCGTTCCACACCACCTGAGGTGATAGTGGAAGTGCTGGAGCCCTATGTCCG CCTTACTGCCAACGTCCGTATCATCAAGAACAGAACAGGCCCTATGGGCCATACCTATGGCTTTATTGACCT CGACTCCCATGCGGAAGCTCTTCGTGTGGTGAAGATCTTACAGAACCTTGATCCGCCATTTAGCATTGATGGGAA GATGGTAGCTGTAAACCTGGCCACTGGAAAACGAAGAATGATTCTGGGGGACCATTCTGACCACATGCATTACTA AGGACAACAGTCATCATGTGACTGCTACATATATGATTCTGCTACTGGCTACTATTATGACCCCTTGGCAGGAAC TTATTATGACCCCAATACCCAGCAAGAAGTCTATGTGCCCCAGGATCCTGGATTACCTGAGGAAGAAGAGATCAA GGAAAAAAACCCACCAGTCAAGGAAAGTCAAGTAGCAAGAAGGAAATGTCTAAAAGAGATGGCAAGGAGAAAAA AGACAGAGGAGTGACGAGGTTTCAGGAAAATGCCAGTGAAGGGCAAGGCCCCTGCAGAAGACGTCTTTAAGAAGCC CCTGCCTCCTACTGTGAAGAAGGAAGAGTCCCCCTCCACCTAAAGTGGTAAACCCACTGATCGGCCTCTTGGG TTGTCTGCTTTGCAGAAGGCAGTTTCCCAATAAAGAAGTTCTGATCAAACACCAGCAGCTGTCAGACCTGCACAA GCAAAACCTGGAAATCCACCGGAAGATAAAACAGTCTGAGCAGGAGCTAGCCTATCTGGAAAGGAGAACGAGA GGGAAAGTTTAAAGGAAGAGGAAATGATCGCAGGGAAAAGCTCCAGTCTTTTGACTCTCCAGAAAGGAAACGGAT TAAGTACTCCAGGGAAACTGACAGTGATCGTAAACTTGTTGATAAAGAAGATATCGACACTAGCAGCAAAGGAGG CTGTGTCCAACAGGCTACTGGCTGGAGGAAAGGGACAGGCCTGGGATATGGCCATCCTGGATTGGCTTCATCAGA GGAGGCTGAAGGCCGGATGAGGGCCCCAGTGTTGGAGCCTCAGGAAGAACCAGCAAAAAGACAGTCCAACGAGAC TTATCGAGATGCTGTTCGAAGAGTCATGTTTGCTCGATATAAAGAACTCGAT**TAA**GAAAGGAGACAAGTTCCATG GGATACAACCTCCCTCTTGTTTTGTCTCTCCTTTTCTTTTGTTACTGTTCTTGCTGCTAGAACTTTTTTAA

## 452/2825 FIGURE 413B

## 453/2825 FIGURE 414

MWGDSRPANRTGPFRGSQEERFAPGWNRDYPPPPLKSHAQERHSGNFPGRDSLPFDFQGHSGPPFANVEEHSFSY
GARDGPHGDYRGGEGPGHDFRGGDFSSSDFQSRDSSQLDFRGRDIHSGDFRDREGPPMDYRGGDGTSMDYRGREA
PHMNYRDRDAHAVDFRGRDAPPSDFRGRGTYDLDFRGRDGSHADFRGRDLSDLDFRAREQSRSDFRNRDVSDLDF
RDKDGTQVDFRGRGSGTTDLDFRDRDTPHSDFRGRHRSRTDQDFRGREMGSCMEFKDREMPPVDPNILDYIQPST
QDREHSGMNVNRREESTHDHTIERPAFGIQKGEFEHSETREGETQGVAFEHESPADFQNSQSPVQDQDKSQLSGR
EEQSSDAGLFKEEGGLDFLGRQDTDYRSMEYRDVDHRLPGSQMFGYGQSKSFPEGKTARDAQRDLQDQDYRTGPS
EEKPSRLIRLSGVPEDATKEEILNAFRTPDGMPVKNLQLKEYNTGYDYGYVCVEFSLLEDAIGCMEANQGTLMIQ
DKEVTLEYVSSLDFWYCKRCKANIGGHRSSCSFCKNPREVTEAKQELITYPQPQKTSIPAPLEKQPNQPLRPADK
EPEPRKREEGQESRLGHQKREAERYLPPSRREGPTFRRDRERESWSGETRQDGESKTIMLKRIYRSTPPEVIVEV
LEPYVRLTTANVRIIKNRTGPMGHTYGFIDLDSHAEALRVVKILQNLDPPFSIDGKMVAVNLATGKRRNDSGDHS
DHMHYYQGKKYFRDRRGGGRNSDWSSDTNRQGQQSSSDCYIYDSATGYYYDPLAGTYYDPNTQQEVYVPQDPGLP
EEEEIKEKKPTSQGKSSSKKEMSKRDGKEKKDRGVTRFQENASEGKAPAEDVFKKPLPPTVKKEESPPPPKVVNP
LIGLLGEYGGDSDYEEEEEEEQTPPPQPRTAQPQKREEQTKKENEEDKLTDWNKLACLLCRRQFPNKEVLIKHQQ
LSDLHKQNLEIHRKIKQSEQELAYLERREREGKFKGRGNDRREKLQSFDSPERKRIKYSRETDSDRKLVDKEDID
TSSKGGCVQQATGWRKGTGLGYGHPGLASSEEAEGRMRGPSVGASGRTSKRQSNETYRDAVRRVMFARYKELD

## 454/2825 FIGURE 415A

 ${\tt GACAGCCTCTGGGTCCTCGGTACAGTCTCTGCACCTCGCGCCCCAGCAGGTAAACTAACATT \textbf{ATG} GATTTT}$ TCCAAGCTACCCAAAATACTCGATGAAGATAAAGAAAGCACATTTGGTTATGTGCATGGGGTCTCAGGACCTGTG  ${\tt ATTATTCGATTGGAGGGTGACATGGCTACTATTCAGGTGTATGAAGAAACTTCTGGTGTGTCTGTTGGAGATCCT}$ GTACTTCGCACTGGTAAACCCCTCTCTGTAGAGCTTGGTCCTGGCATTATGGGAGCCATTTTTGATGGTATTCAA AGACCTTTGTCGGATATCAGCAGTCAGACCCAAAGCATCTACATCCCCAGAGGAGTAAACGTGTCTGCTCTTAGC GGAATTGTCAGTGAGAACTCGCTTATCAAACACAAAATCATGTTACCCCCACGAAACAGAGGAACTGTAACTTAC ATTGCTCCACCTGGGAATTATGATACCTCTGATGTTGTCTTGGAGCTTGAATTTGAAGGTGTAAAGGAGAAGTTC ACTGGCCAGAGAGTCCTTGATGCCCTTTTTCCGTGTGTCCAGGGAGGAACTACTGCTATCCCTGGAGCCTTTGGC TGTGGAAAGACAGTGATATCACAGTCTCTATCCAAGTATTCTAACAGTGATGTAATCATCTATGTAGGATGTGGT GAAAGAGGAAATGAGATGTCTGAAGTCCTCCGGGACTTCCCAGAGCTCACAATGGAGGTTGATGGTAAGGTAGAG TCAATTATGAAGAGGACAGCTTTGGTAGCCAATACCTCCAATATGCCTGTTGCTGCTAGAGAAGCCTCTATTTAT ACTGGAATCACACTGTCAGAGTACTTCCGTGACATGGGCTATCATGTCAGTATGATGGCTGACTCTACCTCTAGA TGGGCTGAGGCCCTTAGAGAAATCTCTGGTCGTTTAGCTGAAATGCCTGCAGATAGTGGATATCCAGCCTATCTT GGTGCCCGTCTGGCCTCGTTTTATGAACGAGCAGGCAGGGTGAAATGTCTTGGAAATCCTGAAAGAGAAGGGAGT GTCAGCATTGTAGGAGCAGTTTCTCCACCTGGTGGTGATTTTTCTGATCCAGTTACATCTGCCACTCTTGGTATC GTTCAGGTGTTCTGGGGCTTAGATAAGAAACTAGCTCAACGTAAGCATTTCCCCTCTGTCAATTGGCTCATCAGC GCTAAGGAAATTCTGCAGGAAGAAGAAGACCTGGCAGAAATTGTACAGCTTGTGGGAAAGGCTTCTTTGGCAGAA ACAGATAAAATCACTCTGGAGGTAGCAAAACTTATCAAAGATGATTTCCTACAACAAAATGGATATACTCCTTAT GACAGGTTCTGCCCATTCTACAAGACAGTAGGGATGCTGTCCAACATGATTGCATTTTATGATATGGCTCGTAGA GCTGTTGAAACCACTGCCCAGAGTGACAATAAAATCACATGGTCCATTATTCGTGAGCACATGGGAGACATCCTC TATAAACTTTCCTCCATGAAATTCAAGGATCCACTGAAAGATGGTGAGGCCAAAGATCAAAAGCGACTATGCACAA TTTCCTCAGCAAGCTCCTATGTGTATATTTTCCTGAATTTCTCATCTCAAACCCTTTGCTTCTTTATTGTGCAGC TGTTCTAGTGTTGTGAAGGGCCTCCCTCTTCCTTTATCTGAAGTGGTGAATATAGTAAATATACATTCTGGTTAC ACTACTGTAAACTTGTATGTAGGGTGATGACCCTCTTTGTCCTAGGTGTACCCTTTCCTCATCTCTATTAAATTG TAAACAGGACTACTGCATGTACTCTTTTGCAGTGAATTTGGAATGGAAGGCCAGGTTTCTATAACTTTTGAACA GGTACTTTGTGAAATGACTCAATTTCTATTGTGGTAAGCTCATTGGCAGCTTAGCATTTTGCAAAGGAATTGCTT TGCAGGAAATATTTAATTTTCAAAAACATAATGATTAATGTTCCAATTATGCATCACTTCCCCCAGTATAAATCA TGGGGACGGAGTGTCCCTCTTGTTGCCCAGGCTGGAGTGCAATGGCGTGATCTTGGCTCACTGCAGCCTTCGCCT CCCGGGTTCAAGTGATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGATTACAGGCATGCTCCACCATGCCCAGCT AATTTTGTATTTTTAGTAGAAACGGGGTTTCACCATATTGGTCAGGCTGGTCTCGAACTCCAGACCTCAGGTGAT  $\tt CCGCCCACCTCGGCCTCCCAAACTGCTGGGATTACAGGCGTGAGCCACCGCGCCTGGCCAGGGACTATACTCTTT$ GGTCAAACTTAGAGCAACATTGTCTTATTAAAGCATAGTTTATTTCACTAGAAAAAATTTAATATCAAGGACTAT TACATACTTCATTACTAGGAAGTTCTTTTTAAAATGACACTTAAAACAATCACTGAAAACTTGATCCACATCACA CCCTGTTTATTTTCCTTAAACATCTTGGAAGCCTAAGCTTCTGAGAATCATGTGGCAAGTGTGATGGGCAGTAAA ATACCAGAGAAGATGTTTAGTAGCAATTAAAGGCTGTTTGCACCTTTAAGGACCAGCTGGGCTGTAGTGATTCCT GGGGCCAGAGTGGCATTATGTTTTTACAAAATAATGACATATGTCACATGTTTGCATGTTTGCTTGTTTGAA ACTGATAACAGAATCTGGGTCACATGAAAAAAATCATTTTATCCGTCTTTTAAGTATATGTTTAAAATAATAAT TTATGTGTCTGCATATTGCAGAACAGCTCTGAGAGCAACAGTTTCCCATTAACTCTTTCTGACCAATAGTGCTGG

## 455/2825 FIGURE 415B

## 456/2825 FIGURE 416

MDFSKLPKILDEDKESTFGYVHGVSGPVVTACDMAGAAMYELVRVGHSELVGEIIRLEGDMATIQVYEETSGVSV GDPVLRTGKPLSVELGPGIMGAIFDGIQRPLSDISSQTQSIYIPRGVNVSALSRDIKWDFTPCKNLRVGSHITGG DIYGIVSENSLIKHKIMLPPRNRGTVTYIAPPGNYDTSDVVLELEFEGVKEKFTMVQVWPVRQVRPVTEKLPANH PLLTGQRVLDALFPCVQGGTTAIPGAFGCGKTVISQSLSKYSNSDVIIYVGCGERGNEMSEVLRDFPELTMEVDG KVESIMKRTALVANTSNMPVAAREASIYTGITLSEYFRDMGYHVSMMADSTSRWAEALREISGRLAEMPADSGYP AYLGARLASFYERAGRVKCLGNPEREGSVSIVGAVSPPGGDFSDPVTSATLGIVQVFWGLDKKLAQRKHFPSVNW LISYSKYMRALDEYYDKHFTEFVPLRTKAKEILQEEEDLAEIVQLVGKASLAETDKITLEVAKLIKDDFLQQNGY TPYDRFCPFYKTVGMLSNMIAFYDMARRAVETTAQSDNKITWSIIREHMGDILYKLSSMKFKDPLKDGEAKIKSD YAQLLEDMQNAFRSLED

## 457/2825 FIGURE 417

CACGTCGGCTGCTGGGAAGATCTGGATTCTCGTTTCAGGTCACCATCAGAAAAGCTAAGTTTGCTGTATAGTGAG GATCAGGAGATCTGATCCTGATTGCAGAACCTTCCCTGATTACAGAATCTTGGATGATTTCACAAAAGTTCATCT TCATTGCAGATACCTGCCTTTCTTTCTAGGTTGTATCTCCCACTTCACCCTTCTAGACCATCCCAGAAGATCTAT CAACTAGAAGCAACATGCAGAGAAGTAAAATGAGAGGGGCCTCCTCAGGAAAGAAGACAGCTGGTCCACAGCAGA AAAATCTTGAACCAGCTCTCCCAGGAAGATGGGGTGGTCGCTCTGCAGAGAACCCCCCTTCAGGATCCGTGAGGA AGACCAGAAAGAACAAGCAGAAGACTCCTGGAAACGGAGATGGTGGCAGTACCAGCGGAAGCACCTCAGCCCCCTC GGAAGAAAAGGGCCCGGGCAGACCCCACTGTTGAAAGTGAGGAGGCGTTTAAGAATAGAATGGAGGTTAAAGTGA AGATTCCTGAAGAATTAAAACCATGGCTTGTTGAGGACTGGGACTTAGTTACCAGGCAGAAGCAGCTGTTTCAAC TCCCTGCCAAGAAAATGTAGATGCAATTCTGGAGGAGTATGCAAATTGCAAGAAATCGCAGGGAAATGTTGATA ATAAGGAATATGCGGTTAATGAAGTTGTGGCAGGAATAAAAGAATATTTCAATGTGATGTTGGGCACTCAGCTGC TCTACAAATTTGAGAGGCCCCAGTATGCTGAAATCCTCTTGGCTCACCCTGATGCTCCAATGTCCCAGGTTTATG GAGCACCACCTACTGAGATTATTTGTAAGAATTGGAGCAATGTTGGCCTATACGCCCCTTGATGAGAAAAGCC TTGCATTATTGTTGGGCTATTTGCATGATTTCCTAAAATATCTGGCAAAGAATTCTGCATCTCTTTTACTGCCA  $\tt GTGATTACAAAGTGGCTTCTGCTGAGTACCACCGCAAAGCCCTG\underline{\textbf{TGA}}{GCGTCTACAGACAGCTCACCATTTTTGT$ CCTGTATCTGTAAACACTTTTTGTTCTTAGTCTTTTTCTTGTAAAATTGATGTTCTTTAAAATCGTTAATGTATA ACAGGGCTTATGTTTCAGTTTGTTTTCCGTTCTGTTTTAAACAGAAAATAAAAGGAGTGTAAGCTCCTTTTCTCA TTTCAAAGTTGCTACCAGTGTATGCAGTAATTAGAACAAAGAAGAACATTCAGTAGAACATTTTATTGCCTAGT TGACAACATTGCTTGAATGCTGGTGGTTCCTATCCCTTTGACACTACACAATTTTCTAATATGTGTTAATGCTAT GTGACAAAACGCCCTGATTCCTAGTGCCAAAGGTTCAACTTAATGTATATACCTGAAAACCCATGCATTTGTGCT CTTTTTTTTTTTTTTTTGGTGCTTGAAGTAAAACAGCCCATCCTCTGCAAGTCCATCTATGTTGTTCTTAGGCATTCT ATCTTTGCTCAAATTGTTGAAGGATGGTGATTTGTTTCATGGTTTTTGTATTTGAGTCTAATGCACGTTCTAACA TGATAGAGGCAATGCATTATTGTGTAGCCACGGTTTTCTGGAAAAGTTGATATTTTAGGAATTGTATTTCAGATC TTAAATAAAATTTGTTTCTAAATTTC

# 458/2825 **FIGURE 418**

 ${\tt MSSRKQGSQPRGQQSAEEENFKKPTRSNMQRSKMRGASSGKKTAGPQQKNLEPALPGRWGGRSAENPPSGSVRKTRKNKQKTPGNGDGGSTSEAPQPPRKKRARADPTVESEEAFKNRMEVKVKIPEELKPWLVEDWDLVTRQKQLFQLPAKKNVDAILEEYANCKKSQGNVDNKEYAVNEVVAGIKEYFNVMLGTQLLYKFERPQYAEILLAHPDAPMSQVYGAPHLLRLFVRIGAMLAYTPLDEKSLALLLGYLHDFLKYLAKNSASLFTASDYKVASAEYHRKAL}$ 

## 459/2825 FIGURE 419

GCGACCGAACGCGGCGGCAGCGTTCGCGCGGGGGCCTGCGAAGCGCTGCTCGGGGCCCGCACTGCCCGCGG CCGGCCCGCGAGCGCAGGAGTAAACACCGCCGGAGTCTTGGAGCCGCTGCAGAAGGGAATAAAGAGAGATGCAG GGATTTGTGAGGTTACGGCGCCCCAGCTGCAAGATGCACTAGCCGGCTGAACCCGGGATCGGCTGACTTGTTGGA  $\verb|CTCTTTCTTTCCTCCGTCTTGTTGCATGCAAGAAAATTACAGTCCGCTGCTCGCCCGGCCCTGGGTGCGAGATAT||$ TCAGCCCCGCTCTCTCCCGTGCATTGTGCAACCCAAAGATGAAAGACCGAAGGGGAGAAAGTTAAAGAAATCGCC CACATGCGCTGGATCAGTCCACGGCTTGGGGAAAGGCATCCAGAGAAGGTGGGAGCGGAGAGTTTGAAGTCTTTA  $\tt CCTCTTGACCCAGGCAGGACAC \underline{\textbf{ATG}} \texttt{CAGGCCAAAAAACGCTATTTCATCCTGCTCTCAGCTGGCTCTTGTCTCGC$ CCTTTTGTTTTATTTCGGAGGCTTGCAGTTTAGGGCATCGAGGAGCCACAGCCGGAGAGAAGAACACAGCGGTAG GAATGGCTTGCACCACCCCAGTCCGGATCATTTCTGGCCCCGCTTCCCGGAGCCTCTGCGCCCCTTCGTTCCTTG GGATCAATTGGAAAACGAGGATTCCAGCGTGCACATTTCCCCCCGGCAGAAGCGAGATGCCAACTCCAGCATCTA CAAAGGCAAGAAGTGCCGCATGGAGTCCTGCTTCGATTTCACCCTTTGCAAGAAAAACGGCTTCAAAGTCTACGT ATACCCACAGCAAAAAGGGGGAGAAAATCGCCGAAAGTTACCAAAACATTCTAGCGGCCATCGAGGGCTCCAGGTT CTACACCTCGGACCCCAGCCAGGCGTGCCTCTTTGTCCTGAGTCTGGATACTTTAGACAGAGACCAGTTGTCACC TCAGTATGTGCACAATTTGAGATCCAAAGTGCAGAGTCTCCACTTGTGGAACAATGGTAGGAATCATTTAATTTT TAATTTATATTCCGGCACTTGGCCTGACTACACCGAGGACGTGGGGGTTTGACATCGGCCAGGCGATGCTGGCCAA AGCCAGCATCAGTACTGAAAACTTCCGACCCAACTTTGATGTTTCTATTCCCCTCTTTTCTAAGGATCATCCCAG GACAGGAGGGGAGAGGGGGTTTTTGAAGTTCAACACCCTCCTCTCTCAGGAAGTACATGCTGGTATTCAAGGG GAAGAGGTACCTGACAGGGATAGGATCAGACACCAGGAATGCCTTATATCACGTCCATAACGGGGAGGACGTTGT GCTCCTCACCACCTGCAAGCATGGCAAAGACTGGCAAAAGCACAAGGATTCTCGCTGTGACAGAGACAACACCGA CTCTGAAGTGATTAATTGGAACCAAGCTGCCGTCATAGGCGATGAGAGATTGTTATTACAGATTCCTTCTACAAT CAGGTCTATTCATCAGGATAAAATCCTAGCACTTAGACAGCAGACACAATTCTTGTGGGAGGCTTATTTTCTTC AGTTGAGAAGATTGTATTAACTACACTAGAGATTATTCAGGACAGAATATTCAAGCACATATCACGTAACAGTTT AATATGGAACAACATCCTGGAGGATTGTTCGTACTACCACAGTATTCATCTTATCTGGGAGATTTTCCTTACTA CTATGCTAATTTAGGTTTAAAGCCCCCCTCCAAATTCACTGCAGTCATCCATGCGGTGACCCCCTGGTCTCTCA GTCCCAGCCAGTGTTGAAGCTTCTCGTGGCTGCAGCCAAGTCCCAGTACTGTGCCCAGATCATAGTTCTATGGAA TTGTGACAAGCCCCTACCAGCCAAACACCGCTGGCCTGCCACTGCTGTCGTCGTCGTCGTCATTGAAGGAGAGAG CAAGGTTATGAGCAGCCGTTTTCTGCCCTACGACAACATCATCACAGACGCCGTGCTCAGCCTTGACGAGGACAC GGTGCTTTCAACAACAGAGGTGGATTTCGCCTTCACAGTGTGGCAGAGCTTCCCTGAGAGGATTGTGGGGTACCC GAACATGGTGGACCAATTGTGAGGACATTCTCATGAACTTCCTGGTGTCTGCTGTGACAAATTGCC TCCAATCAAAGTGACCCAGAAGAAGCAGTATAAGGAGACAATGATGGGACAGACTTCTCGGGCTTCCCGTTGGGC TGACCCTGACCACTTTGCCCAGCGACAGAGCTGCATGAATACGTTTGCCAGCTGGTTTGGCTACATGCCGCTGAT CCACTCTCAGATGAGGCTCGACCCCGTCCTCTTTAAAGACCAGGTCTCTATTTTGAGGAAGAAATACCGAGACAT CCAGTGCAGATCCACTCATCAGCAGAGCCAGATTGTGCCAACTATCCAAAAACTTAGATGAGCAGAATGACAAAA AAAAAAAAGGCCAATGAGAACTCAACTCCTGGCTCCTGGGACTGCACCAGACTGCTCCAAACTCACCTCACTGGC AAAA

## 460/2825 FIGURE 420

MQAKKRYFILLSAGSCLALLFYFGGLQFRASRSHSRREEHSGRNGLHHPSPDHFWPRFPEPLRPFVPWDQLENED SSVHISPRQKRDANSSIYKGKKCRMESCFDFTLCKKNGFKVYVYPQQKGEKIAESYQNILAAIEGSRFYTSDPSQ ACLFVLSLDTLDRDQLSPQYVHNLRSKVQSLHLWNNGRNHLIFNLYSGTWPDYTEDVGFDIGQAMLAKASISTEN FRPNFDVSIPLFSKDHPRTGGERGFLKFNTIPPLRKYMLVFKGKRYLTGIGSDTRNALYHVHNGEDVVLLTTCKH GKDWQKHKDSRCDRDNTEYEKYDYREMLHNATFCLVPRGRRLGSFRFLEALQAACVPVMLSNGWELPFSEVINWN QAAVIGDERLLLQIPSTIRSIHQDKILALRQQTQFLWEAYFSSVEKIVLTTLEIIQDRIFKHISRNSLIWNKHPG GLFVLPQYSSYLGDFPYYYANLGLKPPSKFTAVIHAVTPLVSQSQPVLKLLVAAAKSQYCAQIIVLWNCDKPLPA KHRWPATAVPVVVIEGESKVMSSRFLPYDNIITDAVLSLDEDTVLSTTEVDFAFTVWQSFPERIVGYPARSHFWD NSKERWGYTSKWTNDYSMVLTGAAIYHKYYHYLYSHYLPASLKNMVDQLANCEDILMNFLVSAVTKLPPIKVTQK KQYKETMMGQTSRASRWADPDHFAQRQSCMNTFASWFGYMPLIHSQMRLDPVLFKDQVSILRKKYRDIERL

#### 461/2825 FIGURE 421A

TGAGCTGCGAGGCCCGAGAGTCAGAACCTGGGGGAGAGGGATGGTCTCTGCACGGGGGGGAGCCGGAGGAGCCGC CTCTGGGTGGGCAACTTACCCGAGAACGTGCGGGAAGAGAAGATCATCGAGCATTTCAAACGATATGGCCGCGTG GAAAGTGTCAAAATTCTTCCCAAGAGGGGATCTGAAGGAGGAGTGGCTGCCTTTGTGGATTTTGTGGACATCAAA AGTGCACAGAAAGCTCACAACTCGGTCAACAAAATGGGTGACAGAGACCTACGCACGGATTATAATGAACCAGGC ACCATCCCGAGTGCTCGGGGATTGGATGATACAGTTTCCATAGCATCTCGTAGTAGAGAGGTTTCTGGGTTC CTTGATGGGGCTTCAGATAACAGGGAGCGTGCTTATGAACATAGTGCCTATGGACACCATGAACGGGGGACGGGA GGATTTGATCGGACAAGACATTACGATCAGGATTACTATAGAGATCCTCGAGAGCGGACTTTACAACATGGGCTC TATTACGCTTCTCGGAGTCGAAGTCCAAATCGCTTTGATGCTCATGACCCCCGATATGAACCTAGGGCTCGCGAG CAGTTTACACTGCCCAGTGTGGTACACAGGGATATCTACAGGGATGATATTACCCGGGAGGTACGAGGCAGAAGG  $\verb|CTGGCTAGCCAAGCATCTAGACCCACAAGGTCCCCTAGCGGCAGCGGCTCTAGAAGTAGATCCTCCAGTAGTGAT|\\$ TCAATCAGCAGCAGTAGTACCAGCAGTGACAGCAGTGATTCCAGCAGTAGTTCAAGTGATGATTCTCCAGCT CGATCAGTTCAGCCGCAGCCCGCCCCCCCCCCCCCTTCCCAGTTGCTTTCATCTCTGGAAAAAGATGAGCCCCGT AAAAGTTTTGGCATCAAGGTCCAGAATCTTCCAGTACGCTCTACAGATACAAGCCTTAAAGATGGCCTTTTCCAT GAATTTAAGAAATTTGGAAAAGTAACTTCAGTGCAGATACATGGAACTTCAGAAGAGAGGGTATGGTCTGGTATTC TTTCGGCAGCAAGAAAAAGCCTTGACTGCATCAAAAGGAAAACTTTTCTTTGGCATGCAGATTGAA GTAACAGCATGGATAGGTCCAGAAACAGAAAGTGAAATTAAATTTCGCCCCTTGGATGAAAGGATAGATGAATTT CACCCCAAAGCAACAAGAACTCTCTTTATTGGCAACCTTGAAAAAACCACTACTTACCATGACCTTCGCAACATC TTCCAGCGCTTTGGAGAAATTGTGGATATTGACATTAAGAAAGTAAATGGAGTTCCTCAGTATGCGTTTCTGCAA CTGGGTTTTGGAAAGAGCATGCCTACAAACTGCGTGTGGCTAGATGGGCTTTCTTCGAATGTGTCAGATCAGTAT TTAACACGACATTTCTGCCGATATGGGCCTGTGGTAAAGGTGGTGTTTGACCGCTTAAAAGGCATGGCCCTGGTT CTCTACAATGAAATTGAATATGCACAAGCAGCTGTAAAAGAGACCAAAGGGAGGAAAATCGGTGGGAATAAAATT AAGGTGGATTTTGCAAATCGGGAAAGTCAGCTGGCTTTTTATCACTGCATGGAGAAATCTGGTCAAGACATCAGA GACTTTTATGAAATGTTAGCCGAAAGAAGAGAGGGGAACGAAGGGCATCCTACGACTATAACCAAGATCGTACATAT TATGAGAGTGTTCGAACTCCAGGCACTTATCCTGAGGATTCCAGGCGGGACTATCCAGCTCGAGGGAGAGAGTTT TATTCAGAATGGGAAACTTACCAAGGAGACTACTATGAATCACGATACTACGATGATCCTCGGGAATACAGGGAT TACAGGAATGATCCTTATGAACAAGATATTAGGGAATATAGTTACAGGCAAAGGGAACGAGAAAGAGAACGTGAA AGATTTGAGTCTGACCGGGACAGAGACCATGAGAGGAGGCCGATTGAACGAAGTCAAAGTCCTGTTCACTTGCGA CGTCCACAGAGTCCTGGAGCGTCTCCCTCTCAGGCAGAGAGGTTGCCGAGTGATTCTGAGAGGAGGCTTTACAGC CGATCCTCAGACCGGAGTGGAAGCTGTAGCTCACTCTCCCCTCCAAGATATGAGAAACTGGACAAGTCTCGTTTG TTAATACGGAAGGAAAAAGTGGAAAAGGACAAAACTGACAAGCAGAAACGCAAAGGAAAGGTTCACTCCCCTAGT TCTCAGTCTTCAGAAACGGACCAAGAAAATGAGCGAGAGCAAAGCCCTGAAAAGCCCAGGAGTTGTAATAAACTG AGCAGAGAAAGCTGACAAAGAGGGAATAGCGAAAAACCGCCTGGAACTCATGCCTTGCGTGGTTTTTGACTCGA GTGAAAGAGAGAGGGAAAGGTCATTGACCACACTCCTGTGGAAAAGTTGAAAGCCAAGCTTGATAATGACACT GTCAAATCTTCTGCCCTGGACCAGAAACTTCAGGTCTCTCAGACGGAGCCTGCAAAATCTGACTTGTCTAAACTG CTTAAAGCCAGGAAGCACCTCAAGCCTGAGCAGCCTGCAGATGGGGTAAGTGCTGTGGATCTGGAGAAGCTGGAA GCCAGGAAAAGCCAGTTTGCAGATTCCAATTTAAAAGCAGAAAAGCAAAAACCAGAGGTCAAGAAAAGCAGTCCA GAGATGGAGGATGCTCGCGTGCTTTCAAAAAAGCAGCCTGACGTGTCCTCTAGAGAGGTCATTCTGCTGAGGGAA GGAGAGGCTGAAAGAAAGCCTGTGAGGAAAGAAATTCTTAAAAGAGAATCTAAAAAAATCAAACTGGACAGACTT AATACTGTTGCCAGCCCCAAAGACTGTCAGGAGCTTGCCAGTATTTCTGTTGGGTCTGGCTCAAGGCCCAGCTCA GACCTACAAGCAAGACTGGGAGAACTAGCAGGTGAATCTGTGGAAAATCAAGAAGTCCAATCAAAAAAGCCCATT 

#### 462/2825 FIGURE 421B

TATTGCAGTCTTCGTGATGAAACACCTGAACGTAAATCAGGCCAAGAGAAATCACATTCAGTAAATACTGAAGAA AAAATTGGCATTGACATCGATCACACGCAGAGTTACCGAAAACAAATGGAACAGAGTCGTAGGAAACAGCAGATG GAAATGGAAATAGCCAAGTCTGAGAAGTTTGGCAGTCCTAAAAAAGATGTAGATGAATATGAAAGACGTAGCCTC GTCGATTTTGATATCTGCACCAAGCGAGAACGGAATTACAGAAGTTCACGCCAAATCAGCGAAGATTCTGAAAGG ACTGGTGGTTCTCCCAGTGTCCGACATGGTTCCTTCCATGAAGATGAGGATCCCATAGGCTCCCCTAGGCTACTG TCAGTAAAAGGGTCTCCTAAAGTAGATGAAAAAGTCCTCCCCTATTCTAACATAACAGTCAGGGAAGAGTCTTTA AAATTTAATCCTTATGATTCTAGCAGGAGAGAACAGATGGCAGATATGGCCAAAATAAAACTATCTGTCTTGAAT TCTGAAGATGAACTAAATCGTTGGGACTCTCAGATGAAACAGGATGCTGGCAGATTTGATGTGAGTTTCCCAAAC AGCATAATTAAGAGAGATAGCCTTCGAAAAAGGTCTGTACGAGATCTGGAACCTGGTGAGGTGCCTTCTGATTCT GACGAAGATGGTGAACACAAATCCCACTCACCCAGAGCCTCTGCATTATATGAAAGTTCTCGATTGTCTTTTTTA TTGAGGGACAGAGACAAGCTACGTGAGCGAGATGAAAGACTCTCTAGTTCTTTAGAAAGGAACAAATTTTAC TCTTTTGCATTGGATAAGACAATCACCAGACACTAAAGCTTTGCTTGAAAGAGCTAAATCCCTCTCTTCATCT CGTGAAGAAATTGGTCTTTTCTTGATTGGGACTCCCGATTTGCAAATTTTCGAAACAACAAAGATAAAGAAAAG GTTGACTCTGCTCCAAGACCTATTCCATCCTGGTACATGAAAAAGAAGAAAATTAGGACTGATTCAGAAGGGAAA ATGGATGATAAGAAGAGGACCATAAAGAAGAAGAGAGCAAGAGAGGCAGGAATTGTTTGCTTCTCGTTTTTTACAC AGCTCAATCTTTGAACAAGATTCCAAGCGATTGCAGCATCTAGAGAGAAAAGAGGAAGATTCTGACTTCATTTCT GGTAGGATCTATGGGAAGCAGCATCTGAGGGAGCAAACAGCACAACTGATTCCATTCAAGAACCAGTAGTTCTG GAGAAACAGGAAGATACAGAGAATCATCCCAAGACCCCAGAATCTGCTCCTGAGAATAAAGATTCAGAACTGAAA ACTCCACCTTCCGTTGGGCCTCCAAGTGTCACAGTCGTAACTCTAGAATCAGCCCCATCAGCACTAGAGAAGACC ACTGGTGACAAAACGGTAGAGGCGCCTTTGGTAACAGAAGAGAGACTGTGGAGCCAGCTACCGTCTCAGAAGAA GCAAAGCCTGCATCTGAACCTGCTCCTGCCCCTGTGGAACAGCTGGAACAAGTAGACCTGCCCCCAGGAGCAGAC  $\verb|CCCGATAAAGAAGCTGCCATGATGCCTGCGGGTGTTGAGGAAGGTTCATCAGGTGACCAGCCGCCTTATCTGGAT|$ GCCAAGCCTCCAACTCCCGGGGCCTCGTTTTCCCAGGCAGAGAGCAACGTAGATCCAGAGCCTGACAGTACCCAG CCACTTTCAAAACCAGCTCAGAAGTCTGAGGAAGCCAATGAGCCAAAGGCCGAAAAGCCAGACGCCACTGCAGAT GCTGAGCCTGATGCAAACCAGAAAGCCGAAGCTGCTCCTGAGTCTCAGCCCCCAGCTTCTGAAGATTTAGAGGTT GTGAGTATCGTGGAGAAGCCCGTCACAAGGAAGAGTGAGAGGATAGACCGGGAAAAACTCAAGCGGTCCAATTCT  $\verb|CCTCGGGGAGAAGCACAGAAGCTTTTGGAATTGAAGATGGAGGCAGAAGATTACAAGGACTGCTTCTAAAAAC| \\$ TCTGCTGCAGACCTTGAACATCCCGAACCAAGTTTGCCTCTCAGCCGAACAAGGCGCCGGAATGTAAGGAGCGTC TATGCAACCATGGGTGACCATGAAAACCGCTCTCCTGTCAAAGAGCCCGTTGAGCAACCAAGAGTGACCAGAAAG AGATTGGAGCGAGAGCTTCAGGAGGCTGCAGCGGTTCCCACCACCCCTCGGAGGGGAAGGCCTCCAAAGACACGC CGGCGAGCCGATGAAGAGGAGGAGAACGAGGCCAAGGAACCTGCAGAAACACTCAAGCCACCTGAGGGATGGCGG TCGCCAAGGTCCCAGAAAACTGCAGCTGGTGGTGGACCCCAAGGGAAAAAAGGGAAAAAATGAACCGAAGGTGGAT GCTACACGTCCTGAGGCCACCACTGAGGTGGGCCCCCAAATAGGCGTGAAAGAGAGCTCCATGGAACCCAAGGCT GCTGAGGAGGAGGCAGGAGTGAACAGAAACGTGACAGAAAAGATGCTGGCACAGACAAAAACCCCCCTGAAACC TCCAGGTTAGCAGTGGACAAATCTGCAAGTCTGAAAAATGTGGATGCTGCTGTCAGTCCCAGGGGGGGCTGCAGCA CAGGCAGGGGAGAGGGAATCTGGGGTGGTGGCAGTCTCCCCTGAGAAAAGTGAGAGTCCCCAAAAGGAGGATGGT CCGTCCCCAGAAGCCACCCAGTTAGCCAAGCAGATGGAGCTGGAGCAGGCCGTGGAACACATCGCAAAGCTCGCT GAGGCCTCTGCCTGCTGCCTATAAGGCAGATGCACCAGAGGGCCTTGCCCCAGAGGACAGGGACAAGCCTGCA CACCAAGCAAGTGAAACAGAGCTGGCTGCGGCCATCGGCTCCATCAATGACATTTCTGGGGAGCCAGAAAAC CCTTCTGAGGAAGGAATGGAGACAGATGAGGCTGTATCTGGCATCCTGGAAACTGAGGCTGCTACAGAATCTTCT AGGCCTCCAGTCAATGCTCCTGACCCCTCAGCCGGCCCAACAGATACCAAGGAAGCCAGAGGAAATAGCAGTGAA ACCTCACACTCAGTGCCAGAAGCCAAAGGGTCTAAAGAAGTGGAAGTCACTCTTGTTCGGAAAGACAAAGGGCCC

## 463/2825 FIGURE 421C

GAATCCAACCAAGCTCAAGGTGAGAGTCCTGCTGCAAATGAGGGGACAACAGTACAGCACCCCGAAGCCCCACAG GAAGAAAAGCAGAGTGAGAAACCCCATTCCACTCCTCCTCAGTCATGTACTTCTGACCTAAGCAAGATTCCCTCC CCACCTCCCCAGCCAGCACCGGTGGATGAGGAGCCTCAAGCCAGGTTCAGGGTGCATTCCATCATTGAAAGTGAC CCGGTGACCCCACCCAGCGATCCAAGCATCCCCATACCCACACTGCCTTCTGTAACTGCAGCAAAGCTCTCACCT CCTGTCGCCTCTGGGGGGATCCCACACCAGAGCCCCCCTACTAAGGTGACAGAGTGGATCACAAGGCAGGAGGAG CCACGGGCTCAGTCTACTCCAGCTCTTCCCCCAGACACAAAGGCCTCTGATGTTGACACCAGCTCCAGC ACCCTGAGGAAGATTCTCATGGACCCCAAGTATGTGTCTGCCACAAGTGTCACTTCCACAAGTGTCACCACAGCC GAAGAAAAACAGCACCTCCAGTGACAAACAACTCTGAGATACAAGCCTCGGAGGTGCTGGTAGCTGACAAG GAAAAGGTGGCTCCAGTCATTGCTCCCAAAATTACCTCTGTTATTAGCCGGATGCCTGTCAGCATTGACCTGGAA GTGAACGTCTCCCTGGTCCCGGTGAATGCCCTGAAAGGCCCCGTGAAGGGCTCAGTGACCACACTGAAAAGTTTG GTGAGCACCCCTGCTGGGCCCGTGAACGTCCTGAAAGGGCCTGTGAATGTTCTTACGGGGCCCAGTGAATGTTCTC ACCACTCCAGTGAACGCCACGGTGGGCACAGTGAATGCCGCCCCAGGCACAGTCAATGCCGCTGCGAGTGCAGTG . AATGCCACAGCAAGTGCAGTGACCGTCACAGCGGGTGCGGTTACTGCTGCATCTGGTGGTGTAACGGCCACAACA GGCACGGTGACAATGGCAGGGGCAGTGATTGCGCCGTCAACAAAGTGCAAACAGAGAGCGAGTGCTAATGAAAAC AGTCGGTTCCACCCAGGGTCCATGCCTGTGATCGACGATCGTCCGGCAGACGCGGGCTCAGGGGCGGGGCTGCGT GTGAACACTTCTGAAGGGGTTGTGCTCCTGAGTTACTCAGGGCAGAAGACCGAAGGCCCACAGCGGATCAGCGCC CCTGATTCTGTCACAGCATCGCAGCCTCCATCCAAAGGCCCTCAAGCTCCTGCAGGCTATGCGAACGTGGCCACC CATTCCACGTTGGTACTGACCGCCCAGACATATAATGCCTCTCCTGTGATTTCGTCTGTGAAGGCCGATAGGCCA TCCTTGGAGAAGCCCGAGCCCATTCACCTCTCGGTGTCCACGCCTGTCACCCAGGGAGGCACAGTGAAGGTTCTC ACCCAGGGGATCAACACCCCCCTGTGCTGGTTCACAACCAGCTGGTCCTCACCCCAAGCATTGTCACCACAAAC AAGAAGCTTGCTGACCCCGTCACCCTTAAAATCGAGACCAAGGTCCTTCAGCCGGCCAACCTGGGGTCCACGCTC ACGCCCCACCACCCTCCTGCTCTGCCCAGCAAACTGCCTACAGAAGTCAACCATGTCCCCTCGGGGCCCAGCATC CCAGCAGATCGAACTGTCTCCCATTTGGCAGCTGCAAAGCTAGATGCTCATTCTCCTCGACCAAGTGGACCCGGG CCATCCTCATTCCCAAGGGCAAGCCACCCCAGCAGTACTGCATCTACGGCGCTCTCCACCAACGCCACAGTCATG CTGGCTGCAGGCATCCCAGTGCCCCAGTTCATCTCCAGCATCCACCCAGAGCAGTCTGTCATCATGCCACCCCAC AGCATCACCCAGACTGTGTCCCTGAGCCACCTCTCCCAGGGCGAGGTGAGAATGAACACTCCCACGCTGCCCAGT ATCACCTACAGCATCCGGCCAGAAGCGCTTCACTCTCCTCGGGCTCCGCTGCAGCCCCAGCAAATAGAGGTCAGG GCCCCACAGCGTGCCAGCACCCCGCAGCCCCAGCTGGTGTGCCTGCACTGGCCTCCCAGCACCCTCCCGAG GAGGAAGTGCATTATCACCTTCCTGTCGCTCGAGCCACAGCCCCTGTGCAGTCAGAGGTACTAGTCATGCAGTCT GAGTACCGACTGCACCCCTATACTGTGCCACGGGATGTGAGGATCATGGTGCATCCACATGTGACGGCAGTCAGC GCCCCTGCTCCTCATGGTGAGGCCCGTATCCTCACAGTTACCCCCAGTAACCAACTCCAGGGGCTGCCT CTGACCCCTCCTGTGGTGACCCATGGGGTGCAGATTGTGCACTCCAGCGGGGAGCTGTTTCAAGAGTACCGG CTGCCTTCCCGGACCAAGACAGCTGCTCAGGGCCCTCCTCTGAAGGTGAGCCCCTGCAGCCTCCTCAGCCTGTG AGCAAGATGCCTCAAGTGTCCCAGGAGGCAAAGGGGACCCAGACGGGAGTAGAGCAGCCTCGCCTCCCAGCTGGA AAACAAGATTCCTCTCCACACCTGACTTCCCAGAGACCCGTGGATATGGTTCAACTTCTGAAGAAGTACCCCATC GTGTGGCAGGGCCTGCTGGCCCTCAAGAATGACACAGCTGCTGTGCAGCTCCACTTCGTCTCTGGCAACAACGTC

## 464/2825 FIGURE 421D

CTGGCCCATCGGTCCCTGCCCCTTTCTGAAGGAGGGCCCCCACTAAGGATCGCCCAGAGGATGCGGCTGGAGGCA ACGCAGCTGGAAGGGTTGCCCGAAGGATGACGGTGGAGACAGATTACTGTCTGCTGCTGGCTCTGCCCTGTGGC CGTGACCAAGAGGATGTTGTGAGCCAGACCGAGTCCCTCAAGGCTGCCTTCATCACTTACCTGCAGGCCAAGCAG GAGTTCTCTGAGAGTCACCTGTCCCGCCTGGCCCCTGACCTCCTTGCCAGCATCTCCAACATCTCTCCCCACCTC ATGATTGTCATTGCCTCCGTGTGAGCCACTGAGTGGTTATCACCTCAGTGAATCTTCCCAGGGCTCTGCAGTAAA  $\verb|AACAAAGGACAACCCAGCCAAGCAGGAGGAAGAGCTGCCGAAGGGGGACAGACTCCACTGCCAGACGGCCAGCCGT|$  $\verb|TTGCTGTCCTGCCGGCCCAGTCGGCCAGACTTCCTCTAGGAGTGGTGCTGCTACCTTGTATGTTTACATAA||$  $\verb|TTGCACTAAATTAGTGACTTTTGGCCCAGTGAAGACAGGCTGTGACACTCTGGATGTCTTGGTGTGTA|\\$ GACACACTTGCAGACTCTTAACGCAGGAAGGACTTCAAACTTCTGCTGAGACCTTGGGGTCAAGGAACATTTCA  $\tt TTGGTTTTTTTTGTCCACCCCATCTCCCTTGCTCATTTGGATGCGTCACCTTAATTCTCCTGCTGCCACCGTCT$  $\tt TTGATTCACCGGGATGTACAGTTTACAGTTGAAGAGCAAACAGAAAGGTTTTCTCTTGGTGGGATATGCAGAACT$  ${\tt CATACGTTTTTTTAATCTGTGCCAAAAATGTGTTTTCAGAGGAAATCTTATTTCATATTCAGACTTTGTATTG}$  $\verb|CCCACTCATTTGTATAAGTGCGCTTCGGTACAGCACGGGTCCTGCTCCCGCGATGTGGAAGTGTCACACGGCACC| \\$ TGTACAAAAAGACTGGCTAACCCCTCTTCCTATTACCTTGATCTCTTCCCCCAACTTCCTAACACTTATTAATTT ATGAAACTGTTTTTCTCAGCGCAGTTTTGTTTTGTGTGTCCATTGGATTACAAACTTTATTAAAAAATATAAAAC AC

#### 465/2825 FIGURE 422

MVRETRHLWVGNLPENVREEKIIEHFKRYGRVESVKILPKRGSEGGVAAFVDFVDIKSAQKAHNSVNKMGDRDLR TDYNEPGTIPSAARGLDDTVSIASRSREVSGFRGGGGGPAYGPPPSLHAREGRYERRLDGASDNRERAYEHSAYG  $\tt HHERGTGGFDRTRHYDQDYYRDPRERTLQHGLYYASRSRSPNRFDAHDPRYEPRAREQFTLPSVVHRDIYRDDIT$ SSDDSPARSVQSAAVPAPTSQLLSSLEKDEPRKSFGIKVQNLPVRSTDTSLKDGLFHEFKKFGKVTSVQIHGTSE ERYGLVFFRQQEDQEKALTASKGKLFFGMQIEVTAWIGPETESENEFRPLDERIDEFHPKATRTLFIGNLEKTTT YHDLRNIFQRFGEIVDIDIKKVNGVPQYAFLQYCDIASVCKAIKKMDGEYLGNNRLKLGFGKSMPTNCVWLDGLS SNVSDQYLTRHFCRYGPVVKVVFDRLKGMALVLYNEIEYAQAAVKETKGRKIGGNKIKVDFANRESQLAFYHCME KSGQDIRDFYEMLAERREERRASYDYNQDRTYYESVRTPGTYPEDSRRDYPARGREFYSEWETYQGDYYESRYYD  ${\tt DPREYRDYRNDPYEQDIREYSYRQRERERERERESDRDRDHERRPIERSQSPVHLRRPQSPGASPSQAERLPSD}$ SERRLYSRSSDRSGSCSSLSPPRYEKLDKSRLERYTKNEKTDKERTFDPERVERERRLIRKEKVEKDKTDKQKRK GKVHSPSSQSSETDQENEREQSPEKPRSCNKLSREKADKEGIAKNRLELMPCVVLTRVKEKEGKVIDHTPVEKLK  ${\tt AKLDNDTVKSSALDQKLQVSQTEPAKSDLSKLESVRMKVPKEKGLSSHVEVVEKEGRLKARKHLKPEQPADGVSA}$ VDLEKLEARKRRFADSNLKAEKQKPEVKKSSPEMEDARVLSKKQPDVSSREVILLREGEAERKPVRKEILKRESK KIKLDRLNTVASPKDCQELASISVGSGSRPSSDLQARLGELAGESVENQEVQSKKPIPSKPQLKQLQVLDDQGPE  $\verb"REDVRKNYCSLRDETPERKSGQEKSHSVNTEEKIGIDIDHTQSYRKQMEQSRRKQQMEMEIAKSEKFGSPKKDVD"$ EYERRSLVHEVGKPPQDVTDDSPPSKKKRMDHVDFDICTKRERNYRSSRQISEDSERTGGSPSVRHGSFHEDEDP IGSPRLLSVKGSPKVDEKVLPYSNITVREESLKFNPYDSSRREQMADMAKIKLSVLNSEDELNRWDSQMKQDAGR FDVSFPNSIIKRDSLRKRSVRDLEPGEVPSDSDEDGEHKSHSPRASALYESSRLSFLLRDREDKLRERDERLSSS LERNKFYSFALDKTITPDTKALLERAKSLSSSREENWSFLDWDSRFANFRNNKDKEKVDSAPRPIPSWYMKKKKI  ${\tt RTDSEGKMDDKKEDHKEEEQERQELFASRFLHSSIFEQDSKRLQHLERKEEDSDFISGRIYGKQTSEGANSTTDS}$  ${\tt IQEPVVLFHSRFMELTRMQQKEKEKDQKPKEVEKQEDTENHPKTPESAPENKDSELKTPPSVGPPSVTVVTLESA}$ PSALEKTTGDKTVEAPLVTEEKTVEPATVSEEAKPASEPAPAPVEQLEQVDLPPGADPDKEAAMMPAGVEEGSSG DQPPYLDAKPPTPGASFSQAESNVDPEPDSTQPLSKPAQKSEEANEPKAEKPDATADAEPDANQKAEAAPESQPP ASEDLEVDPPVAAKDKKPNKSKRSKTPVQAAAVSIVEKPVTRKSERIDREKLKRSNSPRGEAQKLLELKMEAEKI TRTASKNSAADLEHPEPSLPLSRTRRRNVRSVYATMGDHENRSPVKEPVEQPRVTRKRLERELQEAAAVPTTPRR GRPPKTRRRADEEEENEAKEPAETLKPPEGWRSPRSQKTAAGGGPQGKKG

## 466/2825 FIGURE 423

# 467/2825 FIGURE 424

 ${\tt MAGSGVRQATSTASTFVKPIFSRDMNEAKRRVRELYRAWYREVPNTVHQFQLDITVKMGRDKVREMFMKNAHVTD} \\ {\tt PRVVDLLVIKGKIELEETIKVWKQRTHVMRFFHETEAPRPKDFLSKFYVGHDP}$ 

## 468/2825 FIGURE 425

GTTGCTGCTAAGCGAACGCCCTTTGGAGCTTACGGAGGCCTTCTGAAAGACTTCACTGCTACTGACTTGTCTGAA  $\verb|TTTGCTGCCAAGGCTGCCTTGTCTGCTGGCAAAGTCTCACCTGAAACAGTTGACAGTGTGATTATGGGCAATGTC| \\$ CTGCAGAGTTCTTCAGATGCTATATATTTGGCAAGGCATGTTGGTTTGCGTGTGGGAATCCCAAAGGAGACCCCA GCTCTCACGATTAATAGGCTCTGTGGTTCTGGTTTTCAGTCCATTGTGAATGGATGTCAGGAAATTTGTGTTAAA  ${\tt GAAGCTGAAGTTGTTTATGTGGAGGAACCGAAAGCATGAGCCAAGCTCCCTACTGTGTCAGAAATGTGCGTTTT}$ GGAACCAAGCTTGGATCAGATATCAAGCTGGAAGATTCTTTATGGGTATCATTAACAGATCAGCATGTCCAGCTC CCCATGGCAATGACTGCAGAGAATCTTACTGTAAAACACAAAATAAGCAGAGAAGAATGTGACAAATATGCCCTG CAGTCACAGCAGAGATGGAAAGCTGCTAATGATGCTGGCTACTTTAATGATGAAATGGCACCAATTGAAGTGAAG ACAAAGAAAGGAAAACAGACAATGCAGGTAGACGAGCATGCTCGGCCCCAAACCACCCTGGAACAGTTACAGAAA CTTCCTCCAGTATTCAAGAAAGATGGAACTGTTACTGCAGGGAATGCATCGGGTGTAGCTGATGGTGCTGGAGCT GTTATCATAGCTAGTGAAGATGCTGTTAAGAAACATAACTTCACACCACTGGCAAGAATTGTGGGCTACTTTGTA TCTGGATGTGATCCCTCTATCATGGGTATTGGTCCTGTCCCTGCTATCAGTGGGGCACTGAAGAAAGCAGGACTG AGTCTTAAGGACATGGATTTGGTAGAGGTGAATGAAGCTTTTGCTCCCCAGTACTTGGCTGTTGAGAGGAGTTTG GATCTTGACATAAGTAAAACCAATGTGAATGGAGGAGCCATTGCTTTGGGTCACCCACTGGGAGGATCTGGATCA AGAATTACTGCACACCTGGTTCACGAATTAAGGCGTCGAGGTGGAAAATATGCCGTTGGATCAGCTTGCATTGGA  ${\tt GGTGGCCAAGGTATTGCTGTCATCATTCAGAGCACAGCC}{{\tt TGA}}{\tt AGAGACCAGTGAGCTCACTGTGACCCATCCTTA}$ CTCTACTTGGCCAGGCCACAGTAAAACAAGTGACCTTCAGAGCAGCTGCCACAACTGGCCATGCCCTGCCATTGA AACAGTGATTAAGTTTGATCAAGCCATGGTGACACAAAAATGCATTGATCATGAATAGGAGCCCATGCTAGAAGT ACATTCTCTCAGATTTGAACCAGTGAAATATGATGTATTTCTGAGCTAAAACTCAACTATAGAAGACATTAAAAG CTTAACTTC

## 469/2825 FIGURE 426

MRLLRGVFVVAAKRTPFGAYGGLLKDFTATDLSEFAAKAALSAGKVSPETVDSVIMGNVLQSSSDAIYLARHVGL RVGIPKETPALTINRLCGSGFQSIVNGCQEICVKEAEVVLCGGTESMSQAPYCVRNVRFGTKLGSDIKLEDSLWV SLTDQHVQLPMAMTAENLTVKHKISREECDKYALQSQQRWKAANDAGYFNDEMAPIEVKTKKGKQTMQVDEHARP QTTLEQLQKLPPVFKKDGTVTAGNASGVADGAGAVIIASEDAVKKHNFTPLARIVGYFVSGCDPSIMGIGPVPAI SGALKKAGLSLKDMDLVEVNEAFAPQYLAVERSLDLDISKTNVNGGAIALGHPLGGSGSRITAHLVHELRRRGGK YAVGSACIGGGQGIAVIIQSTA

#### 470/2825 FIGURE 427

CGCGCCAGGGCGAGGGCACCGCCGGCCGGTCGGGCGCGCTGGGCCTGCCCGGGAATCCCGCCGCCCCGCGCGC CCCGCGCCCTGCGGGCCATGGGAGCCGGCCGGCAGGGACGACGCCTGTGAGACCCGCGAGCGGCCTCGGGGA GGCACGAGAAAGTGAATGGCTTGGAGGAAGGCGTGGAGTTCCTGCCAGTCAACAACGTCAAGAAGGTGGAAAAGC ATGGCCCGGGGCGCTGGTGCTGCCAGCCGTGCTGATCGGCCTCCTCTTGGTCTTGCTGGGGATCGGCTTCC TGGTGTGGCATTTGCAGTACCGGGACGTGCGTGTCCAGAAGGTCTTCAATGGCTACATGAGGATCACAAATGAGA ATTTTGTGGATGCCTACGAGAACTCCAACTCCACTGAGTTTGTAAGCCTGGCCAGCAAGGTGAAGGACGCGCTGA AGCTGCTGTACAGCGGAGTCCCATTCCTGGGCCCCTACCACAAGGAGTCGGCTGTGACGGCCTTCAGCGAGGGCA GCGTCATCGCCTACTACTGGTCTGAGTTCAGCATCCCGCAGCACCTGGTGGAGGAGGCCGAGCGCGTCATGGCCG  ${\tt AGGAGCGCGTAGTCATGCTGCCCCGGGGGGGGCGCTCCCTGAAGTCCTTTGTGGTCACCTCAGTGGTGGCTTTCC}$ CCACGGACTCCAAAACAGTACAGAGGACCCAGGACAACAGCTGCAGCTTTGGCCTGCACGCCCGCGGTGTGGAGC TGATGCGCTTCACCACGCCCGGCTTCCCTGACAGCCCCTACCCCGCTCATGCCCGCTGCCAGTGGGCCCTGCGGG GGGACGCCGACTCAGTGCTGAGCCTCACCTTCCGCAGCTTTGACCTTGCGTCCTGCGACGAGCGCGGCAGCGACC CCTACAACCTGACCTTCCACTCCCAGAACGTCCTGCTCATCACACTGATAACCAACACTGAGCGGCGGCATC CCGGCTTTGAGGCCACCTTCTTCCAGCTGCCTAGGATGAGCAGCTGTGGAGGCCGCTTACGTAAAGCCCAGGGGA CATTCAACAGCCCCTACTACCCAGGCCACTACCCAACATTGACTGCACATGGAACATTGAGGTGCCCAACA ACCAGCATGTGAAGGTGCGCTTCAAATTCTTCTACCTGCTGGAGCCCGGCGTGCCTGCGGGCACCTGCCCCAAGG ACTACGTGGAGATCAATGGGGAGAAATACTGCGGAGAGAGGGTCCCAGTTCGTCGTCACCAGCAACAAGAACAAGA TCACAGTTCGCTTCCACTCAGATCAGTCCTACACCGACACCGGCTTCTTAGCTGAATACCTCTCCTACGACTCCA GTGACCCATGCCCGGGGCAGTTCACGTGCCGCACGGGGCGGTGTATCCGGAAGGAGCTGCGCTGTGATGGCTGGG CCGACTGCACCGACCACGGTGAGCTCAACTGCAGTTGCGACGCCGGCCACCAGTTCACGTGCAAGAACAAGT  ${\tt TCTGCAAGCCCCTCTTCTGGGTCTGCGACAGTGTGAACGACTGCGGAGACAACAGCGACGAGCAGGGGTGCAGTT}$ GTCCGGCCCAGACCTTCAGGTGTTCCAATGGGAAGTGCCTCTCGAAAAGCCAGCAGTGCAATGGGAAGGACGACT GTGGGGACGGGTCCGACGAGGCCTCCTGCCCCAAGGTGAACGTCGTCACTTGTACCAAACACACCTACCGCTGCC AGGACTGCGACTGTGGGCTGCGGTCATTCACGAGACAGGCTCGTGTTGTTGGGGGCACGGATGCGGATGAGGGCC AGTGGCCCTGGCAGGTAAGCCTGCATGCTCTGGGGCCAGGGCCACATCTGCGGTGCTTCCCTCATCTCTCCCAACT GGCTGGTCTCTGCCGCACACTGCTACATCGATGACAGAGGATTCAGGTACTCAGACCCCACGCAGTGGACGGCCT TCCTGGGCTTGCACGACCAGAGCCAGCGCAGCGCCCCTGGGGTGCAGGAGCGCAGGCTCAAGCGCATCATCTCCC ACCCCTTCTTCAATGACTTCACCTTCGACTATGACATCGCGCTGCTGGAGCTGGAGAAACCGGCAGAGTACAGCT CCATGGTGCGGCCCATCTGCCGGACGCCTCCCATGTCTTCCCTGCCGGCAAGGCCATCTGGGTCACGGGCT GGGGACACCCCAGTATGGAGGCACTGGCGCGCTGATCCTGCAAAAGGGTGAGATCCGCGTCATCAACCAGACCA CCTGCGAGAACCTCCTGCCGCAGCAGATCACGCCGCGCGTGATGTGCGTGGGCTTCCTCAGCGGCGGCGTGGACT  $\verb|CCTGCCAGGGTGATTCCGGGGGGACCCCTGTCCAGCGTGGAGGCGGATGGGCGGATCTTCCAGGCCGGTGTGGTGA| \\$ GCTGGGGAGACGGCTCAGAGGAACAAGCCAGGCGTGTACACAAGGCTCCCTCTGTTTCGGGACTGGATCA AAGAGAACACTGGGGTA<u>TAG</u>GGGCCGGGGCCACCCAAATGTGTACACCTGCGGGGCCACCCATCGTCCACCCCAG TGTGCACGCCTGCAGGCTGGAGACTGGACCGCTGACTGCACCAGCCCCCAGAACATACACTGTGAACTCAATC TCCAGGGCTCCAAATCTGCCTAGAAAACCTCTCGCTTCCTCAGCCTCCAAAGTGGAGCTGGGAGGTAGAAGGGGA GGACACTGGTGGTTCTACTGACCCAACTGGGGGCAAAGGTTTGAAGACACAGCCTCCCCCGCCAGCCCCAAGCTG AGTGAAGGTGGTGGGGCTGCCGGATCTGGGCTGTGGGGCCCTTTGGGCCACGCTCTTGAGGAAGCCCAGGCTCGGA GGACCCTGGAAAACAGACGGGTCTGAGACTGAAATTGTTTTACCAGCTCCCAGGGTGGACTTCAGTGTGTATT 

#### 471/2825 FIGURE 428

MGSDRARKGGGGPKDFGAGLKYNSRHEKVNGLEEGVEFLPVNNVKKVEKHGPGRWVVLAAVLIGLLLVLLGIGFL
VWHLQYRDVRVQKVFNGYMRITNENFVDAYENSNSTEFVSLASKVKDALKLLYSGVPFLGPYHKESAVTAFSEGS
VIAYYWSEFSIPQHLVEEAERVMAEERVVMLPPRARSLKSFVVTSVVAFPTDSKTVQRTQDNSCSFGLHARGVEL
MRFTTPGFPDSPYPAHARCQWALRGDADSVLSLTFRSFDLASCDERGSDLVTVYNTLSPMEPHALVQLCGTYPPS
YNLTFHSSQNVLLITLITNTERRHPGFEATFFQLPRMSSCGGRLRKAQGTFNSPYYPGHYPPNIDCTWNIEVPNN
QHVKVRFKFFYLLEPGVPAGTCPKDYVEINGEKYCGERSQFVVTSNSNKITVRFHSDQSYTDTGFLAEYLSYDSS
DPCPGQFTCRTGRCIRKELRCDGWADCTDHSDELNCSCDAGHQFTCKNKFCKPLFWVCDSVNDCGDNSDEQGCSC
PAQTFRCSNGKCLSKSQQCNGKDDCGDGSDEASCPKVNVVTCTKHTYRCLNGLCLSKGNPECDGKEDCSDGSDEK
DCDCGLRSFTRQARVVGGTDADEGEWPWQVSLHALGQGHICGASLISPNWLVSAAHCYIDDRGFRYSDPTQWTAF
LGLHDQSQRSAPGVQERRLKRIISHPFFNDFTFDYDIALLELEKPAEYSSMVRPICLPDASHVFPAGKAIWVTGW
GHTQYGGTGALILQKGEIRVINQTTCENLLPQQITPRMMCVGFLSGGVDSCQGDSGGPLSSVEADGRIFQAGVVS
WGDGCAQRNKPGVYTRLPLFRDWIKENTGV

## 472/2825 FIGURE 429

 ${\tt TAGGAATTTCAAGGTGTAGGTCAGATATT} {\tt TGA}{\tt GAATAGGGGGATTTTTTTGATGTGCCTTAAATTATACCAAAGAT}$ TACTAATTATTCCTCTTTGCCCAAAATACTTGCATCCAAGGTTCTAGTCTCTGTTGCTGTTGCTGGTCTTTAGCCC CACTGCTTGCACTGATGTCCCTCCTTTTCACGGAGACCTATCTGAGGTACAGGATGGGGCTGGCACCAGATGATG TCCCACCACAGTCCCTCACCTCCACATGACAGAACCAATTTACACTCAACCATGACCTCACCCTCCT  ${\tt TGGTTTCTCCCTCGATCTGTGGCCCTTTTTGGATGTATTCTTATCTAACAACACAATCCGGAAAGACTGAATTGA}$ ATATTTATACTAATGGTTCATATCCTTTATTGCTCAATGATCTAATTAAAGGGATCATTGCCACATTTCATGTTT ATATTTCTACAATTTGTTTAGAAAACATCTCCTGACCATATCAGTAGCTCGTGTTATCTTTTTTTCAACTGCTTC CCAGAGTCCTAAAACAATAGAAATTTTGGATTGAAAAGTTCAGCATAAGGAGTTTGAGTCAGTAAAGGATGGGAT AAAGGAGTCGAGATGATCAATGAAAAGTATCACAAAAAAGAGATTGATCAACAAGAGAAATAAAAAGCCCAAG AGGAAGTGGTAGGGGAAGGAATTTAAGAACAGCAATAAGTAAAACTCTTAAGTAACTCCAAAAAAGAAAATGGTAC ATTTTGCCAAAGACCACTTATACTTGAGAACATGGAAGAATTTGCCTGATACTCTCTTTGGGGAAAAGAGTCTCT CCTCTTTTCCTCAAACCCCAGTACACTCAGCCTCTCTGCCCCACCTTCTCCTGACTTTGTCCTCACTTGCTTCTG CAGTACATTGGAACCTGAATTGAAAGAAAGTCTTCCTTGAATAATTGGAGTTTGTCTTGAGAGGCAAATATAGCC CCAAGAATCACAAGATTCGAGGACCATGTAGGTCTTTTACGTAGCCCAAATCCATAAATTAGTCTCACTTTTTGT ATTTATCGTTTCATATTAAACCCTCTATATCAAATGTTCATCATGATTTTTGTATGATTTTTATAACTATTTTATT  ${\tt CATTITATTAGATTTATTCTAAAATTTTTTAATGGTAAATTCTTAAACTGTGGAAACCACTGAAGGTGCTTATTA}$ ACTGTTCTCCCAGATTTGTACAAGTATTGGATGATTCCTTGAGTTTACAGCTGTACAAATAGTGTGGAAAATAAA CTTTTTTTAAAAAAGA

473/2825 FIGURE 430

GISRCRSDI

## 474/2825 FIGURE 431

TCTCACAGGCCCTTAGAGCATCGCCAGGTGCAGAGCTCCACAGCTCTCTTTCCCAAGGAGTAATCAGAGGGTGAG AACGTGGAGCCTGGTGGACAGGTGAAAGCACTGGGATCTTTCTGCCCAGAAAGGGGGAAAGTTGCACATTTATATC  $\tt CTAGAGGGAAGCGACAGCAGTGCTTCTCCCTGTGCTGAGGTACAGGAGCC\textbf{ATG} TGGCTAGAAATCCTCCTCACTT$ CAGTGCTGGGCTTTGCCATCTACTGGTTCATCTCCCGGGACAAAGAGGAAACTTTGCCACTTGAAGATGGGTGGT GGGGGCCAGGCACGAGCCAGCCAGGGAGGACGACACCCCTTTCAAGGTGGAAACGTCAGATGAGG AGATCCACGACTTACACCAGAGGATCGATAAGTTCCGTTTCACCCCACCTTTGGAGGACAGCTGCTTCCACTATG AGCTGCCCGCAGGCCATACCCCGAAGCCCTTGCTGATGGTGCACGGCTGGCCCGGCTCTTTCTACGAGTTTTATA AGATCATCCCACTCCTGACTGACCCCAAGAACCATGGCCTGAGCGATGAGCACGTTTTTGAAGTCATCTGCCCTT  $\verb|CCATCCCTGGCTATGGCTTCTCAGAGGCATCCTCCAAGAAGGGGTTCAACTCGGTGGCCACCGCCAGGATCTTT|\\$ ACAAGCTGATGCTGCGGCTTGCGGATTCTACATTCAAGGAGGGGACTGGGGGTCCCTGATCTGCACTA ATATGGCCCAGCTGGTGCCCAGCCACGTGAAAGGCCTGCACTTGAACATGGCTTTTGGTTTTAAGCAACTTCTCTA CCCTGACCCTCCTGGGACAGCGTTTCGGGAGGTTTCTTGGCCTCACTGAGAGGGATGTGGAGCTGCTGTACC CCGTCAAGGAGAAGGTATTCTACAGCCTGATGAGGGAGAGCGGCTACATGCACATCCAGTGCACCAAGCCTGACA CCGTAGGCTCTGCTCTGAATGACTCTCCTGTGGGTCTGGCTGCCTATATTCTAGAGAAGTTTTCCACCTGGACCA ATACGGAATTCCGATACCTGGAGGATGGAGGCCTGGAAAGGAAGTTCTCCCTGGACGACCTGCTGACCAACGTCA TGCTCTACTGGACAACAGGCACCATCATCTCCTCCCAGCGCTTCTACAAGGAGAACCTGGGACAGGGCTGGATGA CCCAGAAGCATGAGCGGATGAAGGTCTATGTGCCCACTGGCTTCTCTGCCTTTCCCTTTTGAGCTATTGCACACGC CTGAAAAGTGGGTGAGGTTCAAGTACCCAAAGCTCATCTCCTATTCCTACATGGTTCGTGGGGGCCACTTTGCGG CCTTTGAGGAGCCGGAGCTGCTCGCCCAGGACATCCGCAAGTTCCTGTCGGTGCTGGAGCGGCAA**TGA**CCCACCC AATGAGTTTGCCTCCGTCCCCATGCTGGGAGCCCACGCTCACCCCTCACCCCTCCAAGCTCACTCCCCA ACCCCCAACTCCGTGTGGTAAGCAACATGGCTTTGATGATAAACGACTTTACTCTA

## 475/2825 FIGURE 432

MWLEILLTSVLGFAIYWFISRDKEETLPLEDGWWGPGTRSAAREDDSIRPFKVETSDEEIHDLHQRIDKFRFTPP LEDSCFHYGFNSNYLKKVISYWRNEFDWKKQVEILNRYPHFKTKIEGLDIHFIHVKPPQLPAGHTPKPLLMVHGW PGSFYEFYKIIPLLTDPKNHGLSDEHVFEVICPSIPGYGFSEASSKKGFNSVATARIFYKLMLRLGFQEFYIQGG DWGSLICTNMAQLVPSHVKGLHLNMALVLSNFSTLTLLLGQRFGRFLGLTERDVELLYPVKEKVFYSLMRESGYM HIQCTKPDTVGSALNDSPVGLAAYILEKFSTWTNTEFRYLEDGGLERKFSLDDLLTNVMLYWTTGTIISSQRFYK ENLGQGWMTQKHERMKVYVPTGFSAFPFELLHTPEKWVRFKYPKLISYSYMVRGGHFAAFEEPELLAQDIRKFLS VLERQ

## 476/2825 FIGURE 433

TCCGCTATCCAGAACCTCCACTCTTTCGACCCCTTTGCTGATGCAAGTAAGGGTGATGACCTGCTTCCTGCTGGC ACTGAGGATTATATCCATATAAGAATTCAACAGAGAAACGGCAGGAAGACCCTTACTACTGTCCAAGGGATCGCT GATGATTACGATAAAAAGAAACTAGTGAAGGCGTTTAAGAAAAAGTTTGCCTGCAATGGTACTGTAATTGAGCAT CCGGAATATGGAGAAGTAATTCAGCTACAGGGTGACCAACGCAAGAACATATGCCAGTTCCTCGTAGAGATTGGA  $\tt CTGGCTAAGGACGATCAGCTGAAGGTTCATGGGTTT{\color{red}{TAA}}{\color{blue}{GTGCTGTGGGCTCACTGAAGCTTAAGTGAGGATTTC}$ CTTGCAATGAGTAGAATTTCCCTTCTCTCCCTTGTCACAGGTTTAAAAACCTCACAGCTTGTATAATGTAACCAT TTGGGGTCCGCTTTTAACTTGGACTAGTGTAACTCCTTCATGCAATAAACTGAAAAGAGCCATGCTGTCTAGTCT TGAAGTCCCTCATTTAAACAGAGGTCAAGCAATAGGCGCCTGGCAGTGTCAAGCCTGAAACCAAGCAATACCGTC ATGTTTCAGCCAAGCCCAGAGCCCTAAGATTACAAACAACTATGGCCGGAACCTCCTCAGCTCTCCCTCTGCAGA GTTCCCTACCCTAAGAGAATGTTACCACCTGAACAGTCCTCGGTGAATCTGAGAGGAGGAGGATGGGGTAAGGCAG AAGCACCAGCTGTACTAGAAGGGAGCTTTTGGTGGTAGATCCCCTGGTGTCTCCAACCTGACTAGGTGGACA GAGCTCAAAGAGGCCCTCTTACCGCTAGCGAGGTGATAGGACATCTGGCTTGCCACAAAGGTCTGTTCGACCAGA CATATCCTAGCTAAGGGATGTCCAAACATCAGAATGTGAGGCCAACCTTCTATCAGAGTTAAACTTTTGACAAGG TGCCATGTGAAGTAACAAACTGGTTTTTGGTTTTTTTTTCCCCTTCAGTTTTAATGTTATGTGTAATGTATTTAA 

# 477/2825 FIGURE 434

 ${\tt MSAIQNLHSFDPFADASKGDDLLPAGTEDYIHIRIQQRNGRKTLTTVQGIADDYDKKKLVKAFKKKFACNGTVIE}\\ {\tt HPEYGEVIQLQGDQRKNICQFLVEIGLAKDDQLKVHGF}$ 

#### 478/2825 FIGURE 435A

GAGGAAGCGATTCTGGGGTTTCTGTTGAACGGTTCTTGTCCGCGAAGATGCGCTTCGGCCTCTGTCAGGGGAC GTTGGGAGGAGTTCACAGATCCGCTCAGCTTTATCGGCCGCATCCGGCCTTTGGCGGAGAAAACCGGCATCTGCA AAATTCGGCCGCCCAAGGACTGGCAGCCTCCATTTGCCTGTGAAGTAAAAAGCTTTCGTTTCACTCCAAGAGTCC AGCGCCTGAATGAACTTGAGGCAATGACCAGAGTGAGATTTGGATTTCTTGGATCAACTAGCAAAATTTTTGGGAAC TTCAAGGATCTACTCTGAAGATCCCTGTGGTAGAGAGAAAAATCCTGGATCTGTATGCTTTGAGCAAGATTGTTG CCAGCAAAGGAGGTTTTGAAATGGTCACCAAAGAGAAGAAATGGTCTAAAGTGGGTAGTCGCTTGGGATATCTGC CAGGAAAAGGAACTGGGTCTCTTTTGAAGTCACATTATGAAAGAATTCTCTACCCATATGAGCTTTTCCAGTCTG GTGTGAGCCTTATGGGTGTGCAGATGCCTAATTTAGATCTTAAAGAAAAAGTGGAGCCTGAGGTTCTCAGCACTG ATACCCAAACTTCCCCAGAGCCAGGCACAAGGATGAACATTCTGCCGAAGAGAACAAGACGTGTGAAGACTCAGT CAGAATCTGGAGATGTGAGAAACACGGAACTGAAGAAACTTCAGATTTTTGGGGCCTGGGCCCAAGGTTGTGG GCTTGGCAATGGGAACAAAAGATAAAGAAGATGAGGTCACCCGAAGACGAAAAGTTACCAACAGGTCAGACGCAT TTAACATGCAAATGAGACAACGGAAAGGCACTCTCTCTGTTAACTTTGTTGATCTCTATGTTTGTATGTTTTGTG CTCCACTACCTGATGTGCCCAAAGGAGACTGGAGGTGTCCTAAATGTGTCGCCGAGGAATGTAGCAAACCTCGAG AAGCCTTTGGATTTGAACAAGCTGTACGAGAGTATACACTTCAGAGCTTTGGAGAGATGGCAGATAATTTTAAGT CTGATTATTTTAATATGCCAGTCCATATGGTTCCCACAGAACTAGTAGAAAAGGAATTTTGGCGGCTGGTAAGCA GCATTGAAGAAGATGTTATTGTGGAATATGGAGCAGATATCTCCTCAAAAGACTTTGGAAGTGGATTTCCGGTGA AGGATGGGCGGAGAAGATTCTGCCAGAAGAAGAGGGAATATGCACTTTCTGGTTGGAATTTGAATAACATGCCTG TCCTGGAACAGTCTGTTCTTGCACATATTAATGTGGACATCTCTGGTATGAAAGTGCCGTGGCTCTATGTGGGAA TGTGCTTCTTCTTTTTGCTGGCACATTGAGGATCACTGGAGTTATTCCATCAACTACTTGCACTGGGGGGAGC  ${\tt CAAAGACATGGTATGGTGCCATCTCATGCTGCAGAGCAACTGGAGGAGGTGATGAGAGAGCTGGCCCCGAGT}$ TATTTGAATCCCAGCCTGATCTTCTGCATCAGTTAGTTACCATCATGAACCCCAACGTGCTAATGGAGCATGGTG TGCCTGTGTACAGGACCAATCAGTGTGCTGGCGAGTTTGTTGTGACATTTCCTCGTGCCTATCACTCTGGATTTA  ${\tt ACCAGGGCTACAACTTTGCTGAAGCTGTGAACTTCTGTACTGCTGACTGGTTGCCCATTGGACGTCAATGTGTAA}$ ATCATTACCGACGCCTAAGGCGCCACTGTGTCTTTTCACACGAGGAACTAATTTTCAAGATGGCAGCAGATCCAG AATGCTTAGATGTGGGGCTGCCATGGTCTGCAAAGAATTGACTCTCATGACTGAAGAAGAAACACGATTAA GAGAGTCTGTTGTACAGATGGGTGTCCTGATGTCAGAAGAAGAAGTGTTTGAACTTGTTCCTGATGATGAGCGGC TCTACCATCCAACTGATCTGTGCCCCTGCCCCATGCAGAAGAAATGTCTTAGATATCGCTACCCATTAGAAGACC TCCCTTCTCTGCTATATGGTGTAAAAGTCAGGGCACAGTCCTATGACACTTGGGTCAGTCGTGTTACAGAAGCAT TGTCTGCTAACTTCAACCACAAAAAAGATTTGATTGAATTGCGAGTAATGCTGGAAGATGCTGAGGATAGGAAAT ACCCAGAGAATGATCTCTTTCGAAAACTCAGGGATGCTGTAAAAGAAGCTGAGACCTGTGCTTCTGTGGCTCAGC TGCTTCTGAGCAAAAAGCAGAAACACAGACAGACCCAGATAGTGGGAGGACTCGGACCAAACTGACAGTGGAAG AATTGAAGGCCTTTGTCCAACAACTTTTTAGTCTTCCGTGTGTCATCAGCCAAGCTCGGCAAGTAAAGAATCTGC AGATGTTGATAGATATGGGCTCTAGTCTCTATGTGGAACTCCCTGAATTACCACGACTGAAGCAAGAGCTACAAC AGGCTCGGTGGTTGGACGAAGTAAGACTGACCTTATCAGATCCGCAACAAGTCACTTTGGATGTCATGAAGAAGC TGATAGACTCTGGGGTAGGGTTGGCACCCCACCATGCTGTGGAGAAAGCAATGGCTGAACTACAGGAGCTCCTTA CAGTCTCTGAACGATGGGAAGAAAAGGCTAAGGTCTGCCTACAGGCAAGACCGAGGCACAGTGTGGCAAGTTTAG AAAGCATTGTGAATGAAGCCAAGAACATTCCAGCCTTTCTACCCAATGTGTTGTCCTTGAAAGAAGCCTTACAAA AGGCTCGAGAATGGACCGCTAAAGTGGAAGCTATTCAGAGTGGCAGCAATTACGCTTATTTGGAGCAGCTTGAGA GCTTGTCTGCGAAAGGACGCCCTATTCCTGTGCGTCTTGAAGCACTGCCGCAAGTGGAATCACAGGTAGCAGCAG CACGGGCATGGAGAACGGACTGGGCGGACGTTTCTTAAGAAGAATTCTAGCCATACATTGTTACAGGTGCTGA GCCCCGGACCGACATTGGTGTATATGGGAGTGGCAAAAATAGGAGGAAAAAAGTAAAAGAACTAATAGAAAAAG 

## 479/2825 FIGURE 435B

CCAAGATGACAATGGTGGACCGCATAGAAGAAGTAAAATTTTGCATTTGCCGCAAGACAGCCAGTGGGTTTATGC GATCCAGCTGGCAAGCTAAAGAAGTAAAATTCCTTTGCCCTCTTTGTATGCGGTCTCGAAGGCCCAGGCTAGAGA CTATTCTGTCACTCCTGGTATCCCTTCAGAAGTTGCCCGTACGGTTGCCTGAAGGAGAGGCCCTGCAGTGTTTGA  ${\tt CAGAACGTGCTATGAGTTGGCAAGATAGAGCGCGGCAGGCTCTAGCCACAGATGAACTATCCTCTGCCCTGGCCA}$ AACTATCTGTGTTGAGCCAGCGTATGGTGGAACAGGCGGCTCGAGAAAAACTGAAAAGATCATCAGTGCAGAAC TCCAAAAAGCAGCTGCCAATCCAGACTTACAGGGACACTTACCTAGTTTCCAGCAGTCTGCTTTTAACCGGGTGG TGAGCAGTGTGTCATCTTCTCCTCGACAAACAATGGACTATGATGATGAAGAAACAGACTCTGATGAAGACATTC GAGAGACATATGGCTACGACATGAAGGACACAGCCAGTGTGAAGTCCTCTAGTAGTCTTTGAACCCAATCTTTTTT GTGATGAAGAGATTCCCATCAAATCCGAGGAGGTGGTGACCCACATGTGGACAGCACCTTCATTTTGTGCAGAGC ATGCTTATTCTTCTGCTTCTAAGAGTTGTTCTCAAGTATTTTTTTGGGAAAGGTTCTAGCACCCCAAGGAAACAAC TGGAAGAACTTATGATGGTTGGAGATCTCCTGGAAGTATCTCTGGACGAGACTCAACACATATGGCGGATTTTGC AGGCCACACCCCCCCTCTGAAGACAGATTCTTGCATATCATGGAGGATGACAGCATGGAAGAGAAACCACTAA AAGTGAAAGGAAAGGACTCTTCAGAGAAGAAACGGAAACGGAAGCTAGAAAAGGTAGAGCAACTTTTTGGAGAAG CAGCCAAAGTTGAAACTTGTGAAAGAGAGACACTGAAAAGAAAAGAGAGAAAAAGGTGCTGGACATCCCCTCAAAGT ATGACTGGTCAGGAGCAGAGGTCTGATGATGAGAATGCTGTGTGCGCAGAACCAGACTGCCAAAGGCCCTGCA AGGACAAGGGAGTTGTATTTGTAACGAAGAAGAGAGAGATAAAAAATATTAGTTTTAAAAGTGTCCTATGTGACT GCTTTTCTAAAAAGGTAGACTGGGTACAATGTGATGGTGGCTGTGATGAGTGGTTTCATCGGGTTTGTGTGGGTG TATCTCCAGAAATGGCTGAAAATGAAGATTACATCTGTATAAACTGTGCAAAGAAGCAGGGGCCAGTTAGCCCAG TGCTTGGTTAGTTTGGGACATGGGGGGACATGGACCACATTGAGACCTTAGTCATCAAGTAGAGTGGTTTATATC ACTTGGAATGTTGCTTCTAAAGATGAATGGCCTTCAGAGAAAGTCCCCTTAGTGCTGGCTTCCTCTTTGCATGGA CTCTGTGGGTTACATTGCTCTATCAACATATCTATGCAGAGGGTGTCTTCTTTGGTACAACAGCCAATATCTCAT GTCTCCTTTGAGTGTGGTTTACTGCATTAAGGCCAGATGCTTAATTGAGCTCTAGGGTGGCTGGTTAGTATTAAT ACATTGGTGTGCTAACAGGGCATATAGGATGTGGCTTTTGTCCAGCTGATAGTAGTTAGAGGCTTACAACTTAGG  ${\tt AGCAGCACCAACTGAAGGTGCTAATTGCTTGGATCTCCTTCATTAGGATAGTTGGAGAGGGATTGGAGTACCACT}$ TTCTTCCACTGTTACCAGGTACTTAATGCCCTAAAGATACAACTAGGAGTAACAGGGCCAAAGTTATTTCTGTTA GACGTCAAGGAATGGTATCACAGTCTATTGACCTCAGCGATTTGTGCTTGTTTTGTGCTAGAAGAACATCCCAAAT AGGAGAACCTCTCACAAGCTGGGGCAGGTCACCTTATCTTTGTAAGATGAGGATATCATCTAGATCAGAAATCTG ACTAGATTGGATTCTGAGGAGAAGAACCTACTACAAGGCAAGGAGCCGTTTTTTGGCTTTGAAAAGTCTTGCTGT TTCCAGACACCAGGCTTCAGGCATGGCCATGAGCAAGACCAGCAAATAACAGCTTTTTCCCTTGCAGCCCTGACC CCAATGTCTGCTGTTTCCAACACTGGTGATTTCTAACTACGGCCCACAGCAGATGCTGTTGAATAACACCATGGC AAAAA

#### 480/2825 FIGURE 436

MAGVGPGGYAAEFVPPPECPVFEPSWEEFTDPLSFIGRIRPLAEKTGICKIRPPKDWOPPFACEVKSFRFTPRVO RLNELEAMTRVRLDFLDQLAKFWELQGSTLKIPVVERKILDLYALSKIVASKGGFEMVTKEKKWSKVGSRLGYLP GKGTGSLLKSHYERILYPYELFQSGVSLMGVQMPNLDLKEKVEPEVLSTDTQTSPEPGTRMNILPKRTRRVKTQS ESGDVSRNTELKKLQIFGAGPKVVGLAMGTKDKEDEVTRRRKVTNRSDAFNMOMRORKGTLSVNFVDLYVCMFCG RGNNEDKLLLCDGCDDSYHTFCLIPPLPDVPKGDWRCPKCVAEECSKPREAFGFEQAVREYTLQSFGEMADNFKS DYFNMPVHMVPTELVEKEFWRLVSSIEEDVIVEYGADISSKDFGSGFPVKDGRRKILPEEEEYALSGWNLNNMPV LEQSVLAHINVDISGMKVPWLYVGMCFSSFCWHIEDHWSYSINYLHWGEPKTWYGVPSHAAEQLEEVMRELAPEL FESQPDLLHQLVTIMNPNVLMEHGVPVYRTNQCAGEFVVTFPRAYHSGFNQGYNFAEAVNFCTADWLPIGRQCVN HYRRLRRHCVFSHEELIFKMAADPECLDVGLAAMVCKELTLMTEEETRLRESVVQMGVLMSEEEVFELVPDDERQ CSACRTTCFLSALTCSCNPERLVCLYHPTDLCPCPMQKKCLRYRYPLEDLPSLLYGVKVRAQSYDTWVSRVTEAL SANFNHKKDLIELRVMLEDAEDRKYPENDLFRKLRDAVKEAETCASVAQLLLSKKQKHRQSPDSGRTRTKLTVEE LKAFVQQLFSLPCVISQARQVKNLLDDVEEFHERAQEAMMDETPDSSKLQMLIDMGSSLYVELPELPRLKQELQQ ARWLDEVRLTLSDPQQVTLDVMKKLIDSGVGLAPHHAVEKAMAELQELLTVSERWEEKAKVCLQARPRHSVASLE SIVNEAKNIPAFLPNVLSLKEALQKAREWTAKVEAIQSGSNYAYLEQLESLSAKGRPIPVRLEALPQVESQVAAA RAWRERTGRTFLKKNSSHTLLQVLSPRTDIGVYGSGKNRRKKVKELIEKEKEKDLDLEPLSDLEEGLEETRDTAM VVAVFKEREQKEIEAMHSLRAANLAKMTMVDRIEEVKFCICRKTASGFMLQCELCKDWFHNSCVPLPKSSSQKKG SSWQAKEVKFLCPLCMRSRRPRLETILSLLVSLQKLPVRLPEGEALQCLTERAMSWQDRARQALATDELSSALAK LSVLSQRMVEQAAREKTEKIISAELQKAAANPDLQGHLPSFQQSAFNRVVSSVSSSPRQTMDYDDEETDSDEDIR ETYGYDMKDTASVKSSSSLEPNLFCDEEIPIKSEEVVTHMWTAPSFCAEHAYSSASKSCSQVFFGKGSSTPRKQP RKSPLVPRSLEPPVLELSPGAKAQLEELMMVGDLLEVSLDETQHIWRILQATHPPSEDRFLHIMEDDSMEEKPLK VKGKDSSEKKRKRKLEKVEQLFGEGKQKSKELKKMDKPRKKKLKLGADKSKKLNKLAKKLAKEEERKKKKEKAAA AKVELVKESTEKKREKKVLDIPSKYDWSGAEESDDENAVCAEPDCORPCKDKGVVFVTKKREIKNISFKSVLCDC FSKKVDWVQCDGGCDEWFHRVCVGVSPEMAENEDYICINCAKKQGPVSPGPAPPPSFIMSYKLPMEDLKETS

## 481/2825 FIGURE 437A

CTCCTCTAGAGCTTTGGAAGGCTGAATGCACTAAAC**ATG**AAGAGCTTGAAAGCGAAGTTCAGGAAGAGTGACACC AATGAGTGGAACAAGAATGATGACCGGCTACTGCAGGCCGTGGAGAATGCGGAGAAGGTGGCCTCACTG CTCGGCAAGAAGGGGGCCAGTGCCACCAAACACGACAGTGAGGGCAAGACCGCTTTCCATCTTGCTGCTGCAAAA GGACACGTGGAATGCCTCAGGGTCATGATTACACATGGTGTGGATGTGACAGCCCAAGATACTACCGGACACAGC GCCTTACATCTCGCAGCCAAGAACAGCCACCATGAATGCATCAGGAGGCTGCTTCAGTCTAAATGCCCAGCCGAA AGTGTCGACAGCTCTGGGAAAACAGCTTTACATTATGCAGCGGCTCAGGGCTGCCTTCAAGCTGTGCAGATTCTC TGCGAACACAAGAGCCCCATAAACCTCAAAGATTTGGATGGGAATATACCGCTGCTTCTTGCTGTACAAAATGGT CACAGTGAGATCTGTCACTTTCTCCTGGATCATGGAGCAGATGTCAATTCCAGGAACAAAGTGGAAGAACTGCT GTAGATTCTCTTGGATACAATGCCTTACATTATTCCAAACTCTCAGAAAATGCAGGAATTCAAAGCCTTCTATTA TCAAAAATCTCTCAGGATGCTGATTTAAAGACCCCAACAAAACCAAAGCAGCATGACCAAGTCTCTAAAATAAGC TCAGAAAGAGTGGAACTCCAAAAACACGCAAAGCTCCACCCCCCTATCAGTCCTACCCAGTTGAGTGATGTC TCTTCCCCAAGATCAATAACTTCGACTCCACTATCGGGAAAGGAATCGGTATTTTTTTGCTGAACCACCCTTCAAG GCTGAGATCAGTTCTATACGAGAAAACAAAGACAGACTAAGTGACAGTACTACAGGTGCTGATAGCTTATTGGAT AAGGAGTTACAAGATAAATTACAGGCCAAATCACCCCAAGGAGGCGGAAGCAGACCTAAGCTTTGACTCATACCAT TCCACCCAAACTGACTTGGGCCCATCCCTGGGAAAACCTGGTGAAACCTCTCCCCCAGACTCCAAATCATCTCCA TCTGTCTTAATACATTCTTTAGGTAAATCCACTACTGACAATGATGTCAGAATTCAGCAACTGCAAGAGATTTTG CAAGATCTACAGAAGAGATTAGAGAGCTCTGAAGCAGAGAGAAAACAGCTACAGGTCGAACTCCAATCCCGAAGG GCAGAACTGGTATGCTTAAACAACACTGAGATTTCAGAGAACAGCTCTGACCTCAGCCAGAAACTTAAAGAAACT CAGAGCAAATACGAGGAGGCTATGAAAGAAGTCCTTAGTGTGCAGAAGCAGATGAAACTCGGTCTTGTCTCACCT CTGCAGAATGCATTAGAAGAAAGTGAAAGAAAATAAAGAGAAAGTGAGAGAGTTAGAGGAAAAACTGGTAGAGAGG GAGAAAGGTACAGTGATTAAGCCACCTGTGGAAGAGTACGAGGAAATGAAAAGTTCATATTGCTCTGTTATTGAG AATATGAATAAGGAGAAAGCATTTTTGTTTGAGAAATACCAAGAAGCCCCAAGAAGAAATCATGAAATTAAAAGAC ACACTAAAAAGTCAGATGACACAGGAAGCCAGTGATGAAGCTGAGGACATGAAAGAAGCCATGAATAGGATGATA GATGAACTCAATAAACAGGTGAGCGAGCTGTCACAGCTGTACAAAGAAGCCCAGGCTGAGCTGGAGGATTACAGG AAGAGGAAATCTCTAGAGGATGTCACAGCTGAATATATCCATAAAGCAGAGCATGAGAAACTGATGCAATTGACA AACGTGTCCAGGGCTAAAGCAGAAGATGCACTGTCTGAAATGAAGTCTCAGTATTCAAAAGTGTTGAATGAGTTG ACCCAGCTCAAACAACTGGTGGATGCACAAAAAGAGAACTCTGTCTCTATCACAGAACATTTGCAAGTGATAACC GAAGTAGCAAAGCTGGAGAAACACTCTTAGAAGAGAAAGCTGCTATGACTGATGCAATGGTACCTCGGTCTTCC AAAGAGAAGGTCCATTCAGAGGTTGTCCAGATTAGAAGTGAGGTCTCACAGGTGAAAAGAGAAAAAGGAAAATATT CAGACTCTCTTGAAATCCAAAGAGCAAGAAGTAAATGAACTTCTGCAAAAATTCCAGCAAGCTCAGGAAGAACTT GCAGAAATGAAAAGATACGCTGAGAGCTCTTCAAAACTGGAGGAAGATAAAGATAAAAGATAAATGAGATGTCG AAGGAAGTCACCAAATTGAAGGAGGCCTTGAACAGCCTCTCCCAGCTCTCCTACTCAACAAGCTCATCCAAAAAGG CAGAGTCAGCAGCTGGAGGCGCTGCAGCAAGTCAAACAGCTCCAGAACCAGCTGGCGGAATGCAAGAAACAA AAAGTACTGAAGCAAATCCTTACCATGTGTAAAAAACCAGTCTCAAAAGAAGTAAAGTGGATTCCTTGGCAGGACA CTGCCCCTTGTCATCTGTCTTTGTGTTAGATCCAGAGTTGTCGGCAGCCGCTGCCATTGTTCTCATTCGTGGTAT GCACTGTGGCCTAGCGTAGCTTCTTCCCTTTCCAAAGGTTTCTGAGGACTTCTCCCAGGAGAAGACTGCCCGCCT CAGAACTGCTTAGAGACTTCAAACCAGCAGAGGTGAAAGTCCCTGTCATCCCTTCAGATTCCAGAGCTGGGATCA GCCATGCCCAGAGGTCTGGTCCTGATGCTGGCAGGGGGGCCCCCTCCTCCATCCCTGACTGGCTGAGTGGCTTTA TCACCACCGAGTGATGTGCTGAGGCCTCCTGCAGTGAATGCTCCTTTCCATTCCTGTACTCGGGCAGTGCCATTCA CTAGTTATGAAGTATCTGCTTAAAACCCTTCATCATGATATCCTGTGGATTTAAAAAACTCTAATTCCATGTTTTC

## 482/2825 FIGURE 437B

TTCCCATCTGCCTTATATATCTCATCACCCTGCTTATCAATATTCAGTTTGATGAGCACTATTAACTAAAATATG  ${\tt AAACITAAAAACAAAAGCAAGTIGICCITAAAAGTICTITITITAAGTAAATIGITGACATACTGCAAATTITCT}$  $\verb|ATGCAAACTTGCCTCCTGCTGTTATCTGTGAAGCTCAGGAAATCCAAACATTTGTGTTTCAACAAGGGACAGTAA||$  ${\tt ACTGTGTGTTTACAGCCAAAAGAAATGCCTCATAGTTCTTAACCTCAACTTTTGTAGAAGTATTTTTTTCTCTGT}$ AATATTTTTATTGGCTCATAAAGATGTTTTCATATCTGAACTCCTAAATAAGTGAAATTACAGTAGATTATATTA ACAAAATACTTTTTAGGTAGCCATGCTTGAGACTTTTTAAAAATATAACTTTTTCCTTAAAGTTTTCAGCTATAG CAAAAGGTAGTTATGTATGCCAGACCTAATATGAGCTGCCACCAACACCCCTAGAACTTTCAGCCATGGTGTCTT CAGAATTGTAGCGCATTTCTGAATCTAGCAAATCCTCCTTTTACCCGTTGAATGTTTTGAATGCCCTGACTCTAC CAGCGCCCATAAATGATCTCTAGAAGGACTGTTAGTACCAATCTGTTTTTCAACTTTGAAGCTAAAAACCCTGAT TGGCTGTGCTGGATGATATGTTGATCTGTATTGGATAAGGACCAATGACAGCCAAAGCAAAAATGGCTTTAAAGCT TGGTGTTACTTTTCTTAAGTTGTTTAATTATAGTTAAGCAATTTCAAAAATGCTCCAAAGAAATGTGAAAGGACC TTTTGTCACAGCACTTCAGAAAATACACAACAGCCCCTTCTGCCCCGCACAGAAATGCTGCAGAGTATATAAAA ACTCTGGTTAAATATTGAAAAGTTATATGCTGTAGTTTTTAGTATTTTGTCTTTGTAATTTACAGAAGTTATTGG 

## 483/2825 FIGURE 438

MKSLKAKFRKSDTNEWNKNDDRLLQAVENGDAEKVASLLGKKGASATKHDSEGKTAFHLAAAKGHVECLRVMITH GVDVTAQDTTGHSALHLAAKNSHHECIRRLLQSKCPAESVDSSGKTALHYAAAQGCLQAVQILCEHKSPINLKDL DGNIPLLLAVQNGHSEICHFLLDHGADVNSRNKSGRTALMLACEIGSSNAVEALIKKGADLNLVDSLGYNALHYS KLSENAGIQSLLLSKISQDADLKTPTKPKQHDQVSKISSERSGTPKTRKAPPPPISPTQLSDVSSPRSITSTPLS GKESVFFAEPPFKAEISSIRENKDRLSDSTTGADSLLDISSEADQQDLLSLLQAKVASLTLHNKELQDKLQAKSP KEAEADLSFDSYHSTQTDLGPSLGKPGETSPPDSKSSPSVLIHSLGKSTTDNDVRIQQLQEILQDLQKRLESSEA ERKQLQVELQSRRAELVCLNNTEISENSSDLSQKLKETQSKYEEAMKEVLSVQKQMKLGLVSPESMDNYSHFHEL RVTEEEINVLKQDLQNALEESERNKEKVRELEEKLVEREKGTVIKPPVEEYEEMKSSYCSVIENMNKEKAFLFEK YQEAQEEIMKLKDTLKSQMTQEASDEAEDMKEAMNRMIDELNKQVSELSQLYKEAQAELEDYRKRKSLEDVTAEY IHKAEHEKLMQLTNVSRAKAEDALSEMKSQYSKVLNELTQLKQLVDAQKENSVSITEHLQVITTLRTAAKEMEEK ISNLKEHLASKEVEVAKLEKQLLEEKAAMTDAMVPRSSYEKLQSSLESEVSVLASKLKESVKEKEKVHSEVVQIR SEVSQVKREKENIQTLLKSKEQEVNELLQKFQQAQEELAEMKRYAESSSKLEEDKDKKINEMSKEVTKLKEALNS LSQLSYSTSSSKRQSQQLEALQQQVKQLQNQLAECKKQHQEVISVYRMHLLYAVQGQMDEDVQKVLKQILTMCKN QSQKK

## 484/2825 FIGURE 439A

CGGTGGAGACACCCCCGGAGAGGCGGAGGGAAGGGAGGCGAGGCTGCACCTGCATGCTTCCCGCCTCCCACTCC  $\tt CCAGCGCCCCGGACCGTGCAGTTCTCTGCAGGACCAGGCCATGGAGCTCGAAGTCCGGCGGGTCCGACAGGCGT$ TCCTGTCCGGCCGGTCGCGACCTCTGCGGTTTCGGCTGCAGCAGCTGGAGGCCCTGCGGAGGATGGTGCAGGAGC GCGAGAAGGATATCCTGACGGCCATCGCCGCCGACCTGTGCAAGAGTGAATTCAATGTGTACAGTCAGGAAGTCA TTACTGTCCTTGGGGAAATTGATTTTATGCTTGAGAATCTTCCTGAATGGGTTACTGCTAAACCAGTTAAGAAGA ACGTGCTCACCATGCTGGATGAGGCCTATATTCAGCCACAGCCTCTGGGAGTGGTGCTGATAATCGGAGCTTGGA ATTACCCCTTCGTTCTCACCATTCAGCCACTGATAGGAGCCATCGCTGCAGGAAATGCTGTGATTATAAAGCCTT CTGAACTGAGTGAAAATACAGCCAAGATCTTGGCAAAGCTTCTCCCTCAGTATTTAGACCAGGATCTCTATATTG TTATTAATGGTGGTGTTGAGGAAACCACGGAGCTCCTGAAGCAGCGATTTGACCACATTTTCTATACGGGAAACA GTCCATGTTATATTGATAAAGATTGTGACCTGGACATTGTTTGCAGACGCATAACCTGGGGAAAATACATGAATT GTGGCCAAACCTGCATTGCACCCGACTATATTCTCTGTGAAGCATCCCTCCAAAATCAAATTGTATGGAAGATTA AGGAAACAGTGAAGGAATTTTATGGAGAAAATATAAAAGAGTCTCCTGATTATGAAAGGATCATCAATCTTCGTC ATTTTAAGAGGATACTAAGTTTGCTTGAAGGACAAAAGATAGCTTTTGGTGGGGAGACTGATGAGGCCACACGCT ACATAGCCCCAACAGTACTTACCGATGTTGATCCTAAAACCAAGGTGATGCAAGAAGAAATTTTTGGACCAATTC TTCCAATAGTGCCTGTGAAAAATGTAGATGAGGCCATAAATTTCATAAATGAACGTGAAAAGCCTCTGGCTCTTT ACGTCATTATGCACCTCAACTCTTTCCCATTTGGAGGAGTGGGTTCCAGTGGGATGGGAGCTTATCACG GAAAACATAGTTTTGATACTTTTTCTCATCAGCGTCCCTGTTTATTAAAAAAGTTTAAAAAGGAGAGGAGGTGCTAACA AACTCAGATATCCTCCCAACAGCCAGTCAAAGGTGGATTGGGGGGAAATTTTTTCTCTTGAAACGGTTCAACAAAG GAATGATCCTGTTCAACCTCCTAGTGCCTCTACTGAATTATTCCTCTTTTAAATGGTTAATGAACCAATAATTTT TAAATCATACCAAAAATAGTAAGAAAATATGCAAACACTCTGTGATCAAACTTAAAAGTCATTGCCATTCATCAT TAATAAAAGTTGCCATTTCAACTACGTCCCAACATTCCCTAATAGGGTATTCAGGGAACCTGTCTTAAATTGTGC TTATCTAAATCTTGGAACTTTGAGCTAGGGGAGGAGAATGTATTAGACTAAATACAAACTGCGGGGTTGTAAGGG AGTCTCAGAACCTCACTGAATCCTTCACTCCAGTTAATGGCACTGCTCACTTCCTGCCTCTGCTGCCACCATCAC TGTGTGAAGCTTTCAAGAGCTTGGTACTTCCCAGGGCTACCGGCAGTCCTCTGTAGTCCAGAGAGGTGAGATTAG GGGACTCAAGGACTTGACCTCCAGGAGTAGGCCCCTGGTCAGAAGTGCCATCTCACCAGTGGTCTTCATTCTTCC TCATTCATTCTTTATCATCCTGTGTTCTGTTTAGTTGCAACAATCTCTTGTGACTAATGTCACTCAAAGCATCTT GTAACCTAGGGCTTCCTGGAAGTTAGTTGCCAAAGTCATGCAAGCATCACCTGTCATTCTTGTGTTGGAGTTATA GAATTCTACATCTTATAAAACCTAACTGGCATTTAAAAAATACTGTGGCCGGGCGTGGTGGCCTCATGCCTGTAAT CCCAGCACTTTGGGAGGCCGAGGTGGGAGGATTGCTTGAGTCCAGGAATTTGAGACCAGCCTGGACAACACAGTG AGACCICATCTATCAAAAAATAAAAATTAGCTAGATGTGGTGGCATGAGCCTGTGTTCCCAGCTGCTTAGGAG GCTGAAGCAGGAGGATTGATTGAGCCTGCGAGGCCAAGGCTGCAGCAGGCTGTGATTGCACCACTGCACTTCAGC TTGGGCAACAGACCCTGTCTCCGAAACAATAAAAATACTGTAATAAAAGTACTTATAAACATACTAA TCCTCTTCAGGACCCTAAAGTTGCAGGTTAGTAGGTCTTCAAGGACAAATCTGTAAGTTTCTTTTCTGTAGTGC AAGTAAAATTTCACTTTTTGAAACTATAGAGAGATCCCTTTCTGATTAGCCTACAGAACTTAAAGTGAGGGAACC TGGATGGTCTGGACAAGAGCGAGTAAGCCACTGCGGTTGGTCATACTGAAGGGAATTGATGGCAAGAGGATCCC CTGAGCAAGTCAGAAGTTACTCTCATCAGTCGTTCATGGTCACAACCTGAGGTACTCTGCTGAGTGGGCAAGGCT ACGTATAATCAACTTTGTTGAATAATTTGTTCTATTAAGGCTGTCTAAAATGTGATGTCTTCATCATAGTATGAA GTGTTGAAAATTAATAACGAGCCTAGTTTAGGAAAAAGCTGCTTAAAACTGTGGCTCTAAGAGAGTAATCATAAA ATACCTTAGATAAAATTGCACTATGGAATTTTCATTGAGTATGTTTAAATTATTGGCTTGTCTACTAATACATCT GCTTCAAAATGAACATATTTCATAAAATTGGCATCAATTTTAATGACGCTCCTGGTATGGAACCTCAGATATACC

# 485/2825 **FIGURE 439B**

# 486/2825 FIGURE 440

MELEVRRVRQAFLSGRSRPLRFRLQQLEALRRMVQEREKDILTAIAADLCKSEFNVYSQEVITVLGEIDFMLENL PEWVTAKPVKKNVLTMLDEAYIQPQPLGVVLIIGAWNYPFVLTIQPLIGAIAAGNAVIIKPSELSENTAKILAKL LPQYLDQDLYIVINGGVEETTELLKQRFDHIFYTGNTAVGKIVMEAAAKHLTPVTLELGGKSPCYIDKDCDLDIV CRRITWGKYMNCGQTCIAPDYILCEASLQNQIVWKIKETVKEFYGENIKESPDYERIINLRHFKRILSLLEGQKI AFGGETDEATRYIAPTVLTDVDPKTKVMQEEIFGPILPIVPVKNVDEAINFINEREKPLALYVFSHNHKLIKRMI DETSSGGVTGNDVIMHFTLNSFPFGGVGSSGMGAYHGKHSFDTFSHQRPCLLKSLKREGANKLRYPPNSQSKVDW GKFFLLKRFNKEKLGLLLLTFLGIVAAVLVKAEYY

## 487/2825 FIGURE 441

TCGCCGAGTCGATCCGAGAGCTGAGGTCATCGACCCCCAGGACCTCCTAGAAGGCCGATACTTCTCCGGAGCCC TACCAGACGATGAGGATGTAGTGGGGCCCGGGCAGGAATCTGATGACTTTGAGCTGTCTGGCTCTGGAGATCTGG ATGACTTGGAAGACTCCATGATCGGCCCTGAAGTTGTCCATCCCTTGGTGCCTCTAGATAACCATATCCCTGAGA GGGCAGGGTCTGGGAGCCAAGTCCCCACGAACCCAAGAAACTAGAGGAGAATGAGGTTATCCCCAAGAGAATCT CACCCGTTGAAGAGTGAGGATGTGTCCAACAAGGTGTCAATGTCCAGCACTGTGCAGGGCAGCAACATCTTTG AGAGAACGGAGGTCCTGGCAGCTCTGATTGTGGGTGGCATCGTGGGCATCCTCTTTGCCGTCTTCCTGATCCTAC TGCTCATGTACCGTATGAAGAAGAAGGATGAAGGCAGCTATGACCTGGGCAAGAAACCCATCTACAAGAAAGCCC TTTTGAAGGGTGGACATTAGGGTAGGGTGAGGTCAACCTAATACTGACTTGTCAGTATCTCCAGCTCTGATTACC  $\tt TTTGAAGTGTTCAGAAGAGACATTGTCTTCTACTGTTCTGCCAGGTTCTTCTTGAGCTTTGGGCCTCAGTTGCCC$ TGGCAGAAAAATGGATTCAACTTGGCCTTTCTGAAGGCAAGACTGGGATTGGATCACTTCTTAAACTTCCAGTTA AGAATCTAGGTCCGCCTCAAGCCCATACTGACCATGCCTCATCCAGAGCTCCTCTGAAGCCAGGGGGCTAACGG CAGGAAATGGGGGAAGGAACTAGAACCACCTGCACCTTGAGATGTTTCTGTAAATGGGTACTTGTGATCACACTA CGGGAATCTCTGTGGTATATACCTGGGGCCATTCTAGGCTCTTTCAAGTGACTTTTTGGAAATCAACCTTTTTTAT TTGGGGGGGAGATGGGGAAAAGAGCTGAGAGTTTATGCTGAAATGGATTTATAGAATATTTGTAAATCTATTT TAGTGTTTGTTCGTTTTTTAACTGTTCATTCCTTTGTGCAGAGTGTATATCTCTGCCTGGGCAAGAGTGTGGAG GTTTTTTTTTATACTTTAAACCCCGCTCCTCATGGCCTTTGGCCCTTTCTGAAGCTGCTTCCTCTTATAAAATAG CTTTTGCCGAAACATAGTTTTTTTTTAGCAGATCCCAAAATATAATGAAGGGGATGGTGGGGATATTTGTGTCTGT TCTTGCCCTGTTCCTGGTGCCTTCTGATGGGGCTGGAATAGTTGAGGTGGATGGTTCTACCCTTTCTGCCTTCTG TTTGGGACCCAGCTGGTGTTCTTTGGTTTGCTTTCTTCAGGCTCTAGGGCTGTGCTATCCAATACAGTAACCACA TCTAGAGGCGTCACTGTATGTAGTTCATGGCTACTGTACTGACAGCGAGAGCATGTCCATCTGTTGGACAGCACT ATTCTAGAGAACTAAACTGGCTTAACGAGTCACAGCCTCAGCTGTGCTGGGACGACCCTTGTCTCCCTGGGTAGG GAGGCCTGGTGGCTTAGCCTGGGCAGGTCGTGTCTCCTCCTGACCCCAGTGGCTGCGGTGAGGGGAACCACCCTC  $\verb|CCCCAGTGCCCTTCCCCTTGGGCTGTTTGGAGTGAGCACCTCCTCTGTAGGCACCTCTCACACTGTTGTCTGTT|\\$ 

# 488/2825 FIGURE 442

MAPARLFALLLLFVGGVAESIRETEVIDPQDLLEGRYFSGALPDDEDVVGPGQESDDFELSGSGDLDDLEDSMIG PEVVHPLVPLDNHIPERAGSGSQVPTEPKKLEENEVIPKRISPVEESEDVSNKVSMSSTVQGSNIFERTEVLAAL IVGGIVGILFAVFLILLLMYRMKKKDEGSYDLGKKPIYKKAPTNEFYA

## 489/2825 **FIGURE 443A**

CCTGGTCAGCGTCCCGGTCGGGAGTTCTCTCCAGGCGGCACGATGCCGAGGAAACAGTGACCCTGAGCGA ACCAAGCCGGGCGGCAGGTGTGGCTTTGATAGCTGGTGGTGCCACTTCCTGGCCTTGGATGAGCCGTACGCCTCT GTAAACCCAACTTCCTCACCTTTGAAACAGCTGCCTGGTTCAGCATTAATGAAGATTAGTCAGTGACAGGCCTGG TGTGCTGAGTCCGCACATAGAAGAATCAAAAATGTCCAAAATGTAACTGGAGAAAAGTGGGCAACTTTTGGGAG TGACTTTTCCACAGGAACTTCTGCAATGTCCCATCAACCTCTCAGCTGCCTCACTGAAAAGGAGGACAGCCCCAG GCAGATGAAAGAGCTCCTGACCGAGAACCACCAGCTGAAAGAAGCCATGAAGCCTAAATAATCAAGCCATGAAAGG ATCAGAAAGGTCATCTGAGGACCCCACTGATGACTCCAGGCTTCCCAGGCCCGAAGCGGAGCAGGAAAAGGACCA GCTCAGGACCCAGGTGGTGAGGCTACAAGCAGAGAAGGCAGACCTGTTGGGCATCGTGTCTGAACTGCAGCTCAA GCTGAACTCCAGCGGCTCCTCAGAAGATTCCTTTGTTGAAATTAGGATGGCTGAAGGAGAAGCAGAAGGGTCAGT AAAAGAAATCAAGCATAGTCCTGGGCCCACGAGAACAGTCTCCACTGGCACGGCATTGTCTAAATATAGGAGCAG ATCTGCAGATGGGGCCAAGAATTACTTCGAACATGAGGAGTTAACTGTGAGCCAGCTCCTGCTGTGCCTAAGGGA AACAAGTAATCGTTCTGAGATTGAAACCCAGACAGAGGGGGGGCACAGAGAAAGAGAATGATGAAGAGAAAGGCCC GGAGACTGTTGGAAGCGAAGTGGAAGCACTGAACCTCCAGGTGACATCTCTGTTTAAGGAGCTTCAAGAGGCTCA TACAAAACTCAGCGAAGCTGAGCTAATGAAGAAGAGACTTCAAGAAAAGTGTCAGGCCCTTGAAAGGAAAAATTC TGCAATTCCATCAGAGTTGAATGAAAAGCAAGAGCTTGTTTATACTAACAAAAAGTTAGAGCTACAAGTGGAAAG CATGCTATCAGAAATCAAAATGGAACAGGCTAAAACAGAGGATGAAAAGTCCAAATTAACTGTGCTACAGATGAC ACACAACAAGCTTCTTCAAGAACATAATAATGCATTGAAAACAATTGAGGAACTAACAAGAAAAGAGTCAGAAAA AGTGGACAGGGCAGTGCTGAAGGAACTGAGTGAAAAACTGGAACTGGCAGAGAAGGCTCTGGCTTCCAAACAGCT GCAAATGGATGAAATGAAGCAAACCATTGCCAAGCAGGAAGAGGACCTGGAAACCATGACCATCCTCAGGGCTCA GATGGAAGTTTACTGTTCTGATTTTCATGCTGAAAGAGCAGCGAGAGAAAATTCATGAGGAAAAGGAGCAACT GCAGAGTCGTCATGGGGCGAGAACAAGTGACTCTGACCAGCAGGCTTACCTTGTTCAAAGAGGAGCTGAGGACAG GGACTGGCGGCAACAGCGGAATATTCCGATTCATTCCTGCCCCAAGTGTGGAGAGGTTCTGCCTGACATAGACAC  ${\tt GTTACAGATTCACGTGATGGATTGCATCATT} {\color{red}{\textbf{TAA}}} {\color{blue}{\textbf{GTTACAGATTCACCTCCCCAAAACTGTTGGTAAATGTCA}}$ GATTTTTCCTCCAAGAGTTGTGCTTTTGTGTTATTTGTTTTCACTCAAATATTTTGCCTCATTATTCTTGTTTN TGGTGGAGTTCCATCAACTCGCAAAGTAGAATCCTTACCTACTACTCTTCTGATAATAATTTTAATATTTTTTAT TTAGATGGAAGAACCTCACTGATTAATTTTAAGAACCTTTTAGGGATGCAGGAACAATGAAGTGGCCACAGTAT TGTGTGTTGTTGTAGTTTTATTGCAGCCACAATAATTTTACCAAAGTTTTCACATAGGCAGTTAGCCTTTACTT AATATCAAGACAAGTGAAAAAATATTGGCATCGATGAAACCGATAACATTGGCCTCATTGGATTTCTTTACCCAT TTCTAAAAAGAACAGCTGCCTATTTTCTTCCTAGGTTAGGTTATATCTTCATAATCACAAGAATTAGTGATGGCA GCTCATTTGGCTTCATTTAAGCTGTATACCTTAGTCATATATCTTTCATTAGTTCTATGGATATGAGCAGATCCCT TGTTTTGTAAGTCTACAATTTGCCAAGAGTAAATAACACTGGACCAGCTGTAAAAGTAAACAGTGTGTTTATGCA TTGAGATACTAAAGCATTTAAGAAAAAATTAAAAGATCTCTTTTGTTTAAATTTTGTCTTAAGAATCTCTCCTAGA CAAACTTGACTATACACACACACACTTCAATAAGAATCAAAATAGTTCATTATATCTTCATGCCAAGAGAACACA

# 490/2825 FIGURE 443B

## 491/2825 FIGURE 444

MSHQPLSCLTEKEDSPSESTGNGPPHLAHPNLDTFTPEELLQQMKELLTENHQLKEAMKLNNQAMKGRFEELSAW TEKQKEERQFFEIQSKEAKERLMALSHENEKLKEELGKLKGKSERSSEDPTDDSRLPRAEAEQEKDQLRTQVVRL QAEKADLLGIVSELQLKLNSSGSSEDSFVEIRMAEGEAEGSVKEIKHSPGPTRTVSTGTALSKYRSRSADGAKNY FEHEELTVSQLLLCLREGNQKVERLEVALKEAKERVSDFEKKTSNRSEIETQTEGSTEKENDEEKGPETVGSEVE ALNLQVTSLFKELQEAHTKLSEAELMKKRLQEKCQALERKNSAIPSELNEKQELVYTNKKLELQVESMLSEIKME QAKTEDEKSKLTVLQMTHNKLLQEHNNALKTIEELTRKESEKVDRAVLKELSEKLELAEKALASKQLQMDEMKQT IAKQEEDLETMTILRAQMEVYCSDFHAERAAREKIHEEKEQLALQLAVLLKENDAFEDGGRQSLMEMQSRHGART SDSDQQAYLVQRGAEDRDWRQQRNIPIHSCPKCGEVLPDIDTLQIHVMDCII

#### 492/2825 FIGURE 445

GGCACGAGGGGCTGCCCCTCCGCCACCGCCGCCGCCGCCGGCAGGTTCCCTGGTCAGCGTCCCATCCCGGTCG GGAGTTCTCTCCAGGCGGCACGATGCCGAGGAAACAGTGACCCTGAGCGAAGCCAAGCCGGGCGGCAGGAACTTC CCCCCACCTGGCCCAAACCTGGACACGTTTACCCCGGAGGAGCTGCTGCAGCAGATGAAAGAGCTCCTGAC  $\tt CGAGAACCACCAGCTGAAAGAAGCCATGAAGCTAAATAATCAAGCCATGAAAGGGAGATTTGAGGAGCTTTCGGC$ CTGGACAGAGAACAGAAGAAGAACGCCAGTTTTTTGAGATACAGAGCAAAGAAGCAAAAGAGCGTCTAATGGC CTTGAGTCATGAGAATGAGAAATTGAAGGAAGAGCTTGGAAAACTAAAAGGGAAATCAGAAAGGTCATCTGAGGA CCCCACTGATGACTCCAGGCCTTCCCAGGGCCGAAGCGGAGCAGGAAAAGGACCAGCTCAGGACCCAGGTGGTGAG GCTACAAGCAGAAGGCAGACCTGTTGGGCATCGTGTCTGAACTGCAGCTGAACTCCAGCGGCTCCTC AGAAGATTCCTTTGTTGAAATTAGGATGGCTGAAGGAGAGCAGAAGGGTCAGTAAAAGAAATCAAGCATAGTCC TGGGCCCACGAGAACAGTCTCCACTGGCACGGCATTGTCTAAATATAGGAGCAGATCTGCAGATGGGGCCAAGAA TTACTTCGAACATGAGGAGTTAACTGTGAGCCAGCTCCTGCTGTGCCTAAGGGAAGGGAATCAGAAGGTGGAGAG ACTTGAAGTTGCACTCAAGGAGGCCAAAGAAAGAGTTTCAGATTTTGAAAAGAAACAAGTAATCGTTCTGAGAT TGAAACCCAGACAGAGGGGGGGCACAGAGAAAGAGAATGATGAAGAGAAAGGCCCGGAGACTGTTGGAAGCGAAGT GGAAGCACTGAACCTCCAGGTGACATCTCTGTTTAAGGAGGCTTCAAGAGGCTCATACAAAACTCAGCGAAGCTGA GCTAATGAAGAAGACTTCAAGAAAAGTGTCAGGCCCTTGAAAGGAAAAATTCTGCAATTCCATCAGAGTTGAA TGAAAAGCAAGAGCTTGTTTATACTAACAAAAGTTAGAGCTACAAGTGGAAAGCATGCTATCAGAAATCAAAAT GGAACAGGCTAAAACAGAGGATGAAAAGTCCAAATTAACTGTGCTACAGATGACACACAACAAGCTTCTTCAAGA ACATAATAATGCATTGAAAACAATTGAGGAACTAACAAGAAAAGGTCAGAAAAAGTGGACAGGGCAGTGCTGAA GGAACTGAGTGAAAAACTGGAACTGGCAGAGAAGGCTCTGGCTTCCAAACAGCTGCAAATGGATGAAATGAAGCA AACCATTGCCAAGCAGGAAGGACCTGGAAACCATGACCATCCTCAGGGCTCAGATGGAAGTTTACTGTTCTGA TTTTCATGCTGAAAGAGCAGCGAGAGAAAATTCATGAGGAAAAGGAGCAACTGGCATTGCAGCTGGCAGTTCT GCTGAAAGAGATGATGCTTTCGAAGACGGAGGCAGGCAGTCCTTGATGGAGATGCAGAGTCGTCATGGGGCGAG AACAAGTGACTCTGACCAGCAGGCTTACCTTGTTCAAAGAGGGGGCTGAGGACAGGGGACTGGCGGCAACAGCGGAA TATTCCGATTCATTCCTGCCCCAAGTGTGGAGAGGTTCTGCCTGACATAGACACGTTACAGATTCACGTGATGGA  $\verb|TTGCATCATT| \underline{\textbf{TAA}} \\ \texttt{GTGTTGATGTATCACCTCCCCAAAACTGTTGGTAAATGTCAGATTTTTCCTCCAAGAGTTG} \\$ GACCAGCCTGGCCAACATGGCGGAACCCTGTCTCTACCAAAATTACAAAAATTAGCCGAGCATGGTGGCGCATGC CTGTAGTCGCAGCTACTCGCGAGGTTGAGGCAGGAGATTGCTTGAACCCAGGAAGTGGCAGTTGCAGTGAGCCG AA

# 493/2825 FIGURE 446

MSHQPLSCLTEKEDSPSESTGNGPPHLAHPNLDTFTPEELLQQMKELLTENHQLKEAMKLNNQAMKGRFEELSAW TEKQKEERQFFEIQSKEAKERLMALSHENEKLKEELGKLKGKSERSSEDPTDDSRLPRAEAEQEKDQLRTQVVRL QAEKADLLGIVSELQLKLNSSGSSEDSFVEIRMAEGEAEGSVKEIKHSPGSTRTVSTGTALSHYRRRSADGAKNY FEHEELTVSQLLLCLREGNQKVERLEVALKEAKERVSDFEKKTSNRSEIETQTEGSTEKENDEEKGPETVGSEVE ALNLQVTSLFKELQEAHTKLSEAELMKKRLQEKCQALERKNSAIPSELNEKQELVYPNKKLELQVESMLSEIKME QAKTEDEKSKLTVLQMTHNKLLQEHNNALKTIEELTRKESEKVDRAVLKELSEKLELAEKALASKQLQMDEMKQT IAKQEEDLETMTILRAQMEVYCSDFHAERAAREKIHEEKEQLALQLAVLLKENDAFEDGGRQSLMEMQSRHGART SDSDQQAYLVQRGAEDRDWRQQRNIPIHSCPKCGEVLPDIDTLQIHVMDCII

#### 494/2825 FIGURE 447

 $\verb|CCGCCGCTGTCGCCGTCGAGTCGCCATGGAAGTGCAGAAAGAGGCACAGCGCATCATGACCCTGTCGGTGTGGAA| \\$ GATGTATCACTCCCGCATGCAGCGGGTGGCCTGCGGCTGCACCGGAGTCTGCAGCTGTCGCTGGTCATGCGCAG TGACCCTCGCCTGCACCCCCGAGAAGCCGAGTCCACGGCCGAGACAGCGACCCCCGACGGTGAGCACCCGTT TCCGGAGCCAATGGACACGCAGGAGGCGCCGACAGCCGAGGAGACCTCCGCCTGTTGCCCCGCGCCCCCCAA AGTCAGCCGCAAACGACGCAGCAGCCTGAGCGACGGCGGGGGACGTTGGACTGGTCCCGAGCAAGAAAGCCCG TCTGGAAGAAAAGGAAGAGGAGGGAGCGTCATCCGAAGTCGCCGATCGCCTGCAGCCCCTCCGGGCCAAGC GGAGGGCGCCTTTCCCAACCTGGCCCGCGTCCTGCAGAGGCGCTTCTCCGGCCTCCTGAACTGCAGCCCCGCGGC CCCTCCGACGCCCCCCCCGCGTGCGAGGCAAAGCCCGCTTGCCGCCCGGCGGACAGCATGCTCAACGTGCTCGT GCGGGCCGTGGTGGCCTTC**TGA**GGACCCCGAGCGGCGCTGCCGGAGCCCAGAGCGCGCGTCGAACCGTCGGCCCG AGGGCGCAGACCTGAGGCGAGGCCACCCCCCTCCATCCTGGGGGAAGCGCCCGCGAAAACCGTGGAGAGAGCCG TTGTTTTTAAAGATCTCCTCAGGGTCGGACTTCATTTTGTACTGTGGGCTGTGCTGGCCCTTTCAAGGTTTTTCA AGAGTTGGTTTTGCGTTTCCAACCTCGGAGAATTCCAGGCACTCCCCTTCCCCCTCCGCTGACATACTTGTATAA GCGGTCATCGTTGCGTCATGGGGCAGCCTTGGGAGCTTCCTGTCGCCTTGGCTGGGTGTGGGCCTGGAGGAAGG TCCTGGGGCGTGCACTCGCCTGGGCAGTGGGGAGAGTGGCCTGAGTTACTTCACCCCCGGGTGCTGCTGTT AATGTCCCGCGTCTCTGCACCTTCGGGTGGGAGCGGGGACTGATCTACTTTCACATTCTCAAGTTTTTCTCATCT GCATTAGAGGTCCCCAGTAGGTTCCCAGGTTCCAGCGTCCCTCAGACACACGGACACAATCAGCCGAGA AGTTCCTGGTCTGAATCACGAGAATGTGGAGGGGTGGGGGGTGTCAGTGGAAAGGCATAAGGCTGAGACC AGTTGCTGGTGAAACTGGGCCAATCTGGGGAGGGGAACATCCTTGCCAGGGAGTTTCTGAGGGTCTGCTTTGTTT ACCTTTCGTGCGGTGGATTCTTTTTAACTCCGTCTACCTGGCGTTTTGTTAGAAATGTCAGATAGGAAAATAAAA ACCATTIGAGT

# 495/2825 FIGURE 448

 ${\tt MEVQKEAQRIMTLSVWKMYHSRMQRGGLRLHRSLQLSLVMRSARELYLSAKVEALEPEVSLPAALPSDPRLHPPR}\\ {\tt EAESTAETATPDGEHPFPEPMDTQEAPTAEETSACCAPRPAKVSRKRRSSSLSDGGDVGLVPSKKARLEEKEEEE}\\ {\tt GASSEVADRLQPPPGQAEGAFPNLARVLQRRFSGLLNCSPAAPPTAPPACEAKPACRPADSMLNVLVRAVVAF}\\ {\tt CASSEVADRLQPPPGQAEGAFPNLARVLQRRFSGLLNCSPAAPPTAPPACEAKPACRPADSMLNVLVRAVVAF}\\ {\tt CASSEVADRLQPPGAGAFPNLARVLQRRFSGLLNCSPAAPPTAPPACEAKPACRPADSMLNVLVRAVVAF}\\ {\tt CASSEVADRLQPPGAGAFPNLARVLQRRFSGLLNCSPAAPPTAPPACEAKPACRPADSMLNVLVRAVVAF}\\ {\tt CASSEVADRLQPPGAGAFPNLARVLQRRFSGLLNCSPAAPPTAPPACEAKPACRPADSMLNVLVRAVVAF}\\ {\tt CASSEVADRLQPPGAGAFPNLARVLQRRFSGLLNCSPAAPPTAPPACEAKPACRPADSMLNVLVRAVVAF}\\ {\tt CASSEVADRLQPPGAGAFPNLARVLQPGAGAFPNLARVLQPPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPGAGAFPNLARVLQPG$ 

#### 496/2825 FIGURE 449

GCCGGCCGAACCCAGACCCGAGGTTTTAGAAGCAGAGTCAGGCGAAGCTGGGCCAGAACCGCGACCTCCGCAACC TTGAGCGGCATCCGTGGAGTGCGCCTGCGCAGCTACGACCGCAGCAGGAAAGCGCCGCCGGCCAGGCCCAGCTGT GGCCGGACAGGGACTGGAAGAGAGGCGCGGTCGAGTAGGTGTGCACCAGCCCTGGCAACGAGAGCGTCTACCCC GAACTCTGCTGGCCTTGAGGTGGGGAAGCCGGGGGGGGCAGTTGAGGACCCCGCGGAGGCGCGTGACTGGTTGAG  $\tt CGGGCAGCCAGCCTCCGAGCCGGGTGGACACAGGTTTTAAAAC\textbf{ATG} AATCCTACACTCATCCTTGCTGCCTTTT$ GCCTGGGAATTGCCTCAGCTACTCTAACATTTGATCACAGTTTAGAGGCACAGTGGAACGCGAAGGCGATGC ACAACAGATTATACGGCATGAATGAAGAAGGATGGAGGAGGAGCAGTGTGGGAGAAGAACATGAAGATGATTGAAC TGCACAATCAGGAATACAGGGAAGGGAAACACAGCTTCACAATGGCCATGAACGCCTTTGGAGACATGACCAGTG AAGAATTCAGGCAGGTGATGAATGGCTTTCAAAACCGTAAGCCCAGGAAGGGGAAAGTGTTCCAGGAACCTCTGT GTTCTTGTTGGGCTTTTAGTGCTACTGGTGCTCTTGAAGGACAGATGTTCCGGAAAACTGGGAGGCTTATCTCAC TGAGTGAGCAGAATCTGGTAGACTGCTCTGGGCCTCAAGGCAATGAAGGCTGCAATGGTGGCCTAATGGATTATG CTTTCCAGTATGTTCAGGATAATGGAGGCCTGGACTCTGAGGAATCCTATCCATATGAGGCAACAGAAGAATCCT GTAAGTACAATCCCAAGTATTCTGTTGCTAATGACACCGGCTTTGTGGACATCCCTAAGCAGGAGAAGGCCCTGA AAGGCATTTATTTTGAGCCAGACTGTAGCAGTGAAGACATGGATCATGGTGTGTGGTTGGCTACGGATTTG AAAGCACAGAATCAGATAACAATAAATATTGGCTGGTGAAGAACAGCTGGGGTGAAGAATGGGGCATGGGTGGCT CCGCTGTGTCGGATACACTCGAATCATTGAAGATCCGAGTGTGATTTTGAATTCTGTGATATTTTCACACTGGT AAATGTTACCTCTATTTTAATTACTGCTATAAATAGGTTTATATTATTGATTCACTTACTGACTTTGCATTTTCG TTTTTAAAAGGATGTATAAATTTTACCTGTTTAAATAAAATTTAATTTCAAATGTA

## 497/2825 FIGURE 450

MNPTLILAAFCLGIASATLTFDHSLEAQWTKWKAMHNRLYGMNEEGWRRAVWEKNMKMIELHNQEYREGKHSFTM AMNAFGDMTSEEFRQVMNGFQNRKPRKGKVFQEPLFYEAPRSVDWREKGYVTPVKNQGQCGSCWAFSATGALEGQ MFRKTGRLISLSEQNLVDCSGPQGNEGCNGGLMDYAFQYVQDNGGLDSEESYPYEATEESCKYNPKYSVANDTGF VDIPKQEKALMKAVATVGPISVAIDAGHESFLFYKEGIYFEPDCSSEDMDHGVLVVGYGFESTESDNNKYWLVKN SWGEEWGMGGYVKMAKDRRNHCGIASAASYPTV

#### 498/2825 FIGURE 451

 ${\tt AATTCCGCGGAATCATCGGAATCCTTCACC} \underline{{\tt ATC}} {\tt GCATCCAGCCCGGCCCAGCGTCGGCGAGGCAATGATCCTCTC}$ ACCTCCAGCCCTGGCCGAAGCTCCCGGCGTACTGATGCCCTCACCTCCAGCCCTGGCCGTGACCTTCCACCATTT GAGGATGAGTCCGAGGGGCTCCTAGGCACAGAGGGGCCCCTGGAGGAAGAAGAGGATGGAGAGGAGCTCATTGGA GATGGCATGGAAAGGGACTACCGCGCCATCCCAGAGCTGGACGCCTATGAGGCCGAGGGACTGGCTCTGGATGAT GAGGACGTAGAGGAGCTGACGGCCAGTCGAAGGGAGGCAGCAGACGGGCCATGCGGCACGGTGACCGGGAGCTGG CCGGGGCTGGGCGCATGCGCCGTGGGCTCCTGTATGACAGCGATGAGGAGGACGAGGAGGACGCCCTGCCCGCAAGC GCCGCCAGTGGAGCCCGGCAGGAGGACGAGGAGGACGAGATGATTGAGAGCATCGAGAACCTGGAGGAT  $\verb|CTCAAAGGCCACTCTGTGCGCGAGTGGGTGAGCATGGCGGGCCCCCGGCTGGAGATCCACCACCGCTTCAAGAAC| \\$  $\verb|TTCCTGCGCACTCACGTCGACAGCCACGGCCACAACGTCTTCAAGGAGCGCATCAGCGACATGTGCAAAGAGAAC|$  $\verb|CCGGCGGAGCTGCTGCAGATCTTTGATGAGGCTGCCCTGGAGGTGCTACTGGCCATGTACCCCAAGTACGACCGC| \\$ ATCACCAACCACATCCATGTCCGCATCTCCCACCTGCCTCTGGTGGAGGGCTGCGCTGCGCTGAGGCAGCTGCAT  $\tt CTGAACCAGCTGATCCGCACCAGTGGGGTGACCAGCTGCACTGGCGTCCTGCCCCAGCTCAGCATGGTCAAG$ TACAACTGCAACAAGTGCAATTTCGTCCTGGGTCCTTTCTGCCAGTCCCAGAACCAGGAGGTGAAACCAGGCTCC TGTCCTGAGTGCCAGTCGGCCGGCCCCTTTGAGGTCAACATGGAGGAGACCATCTATCAGAACTACCAGCGTATC CGAATCCAGGAGAGTCCAGGCAAAGTGGCGGCTCGGCGGCTGCCCGCTCCAAGGACGCCATTCTCCTCGCAGAT CTGGTGGACAGCTGCAACGCAGGAGACGAGATAGAGCTGACTGGCATCTATCACAACAACTATGATGGCTCCCTC AACACTGCCAATGGCTTCCCTGTCTTTGCCACTGTCATCCTAGCCAACCACGTGGCCAAGAAGGACAACAAGGTT GCTGTAGGGGAACTGACCGATGAAGATGTGAAGATGATCACTAGCCTCTCCAAGGATCAGCAGATCGGAGAGAAG ATCTTTGCCAGCATTGCTCCTTCCATCTATGGTCATGAAGACATCAAGAGAGGCCCTGCTCTGGCCCTGTTCGGA GGGGAGCCCAAAAACCCAGGTGGCAAGCACAAGGTACGTGGTGATATCAACGTGCTCTTGTGCGGAGACCCTGGC ACAGCGAAGTCGCAGTTTCTCAAGTATATTGAGAAAGTGTCCAGCCGAGCCATCTTCACCACTGGCCAGGGGGCG TCGGCTGTGGCCGTCACGGCGTATGTCCAGCGGCACCCTGTCAGCAGGGAGTGGACCTTGGAGGCTGGGGCCCTG GAGGCCATGGAGCAACAGAGCATCTCCATCTCGAAGGCTGGCATCGTCACCTCCCTGCAGGCTCGCTGCACGGTC ATTGCTGCCGCCAACCCCATAGGAGGCGCTACGACCCCTCGCTGACTTTCTCTGAGAACGTGGACCTCACAGAG  $\verb|CCCATCATCTCACGCTTTGACATCCTGTGTGTGGGGGACACCGTGGACCCAGTCCAGGACGAGATGCTGGCC| \\$ CGCTTCGTGGTGGGCAGCCACGTCAGACACCCCCAGCAACAAGGAGGAGGGGGGCTGGCCAATGGCAGCGCT GCTGAGCCCGCCATGCCCAACACGTATGGCGTGGAGCCCCTGCCCCAGGAGGTCCTGAAGAAGTACATCATCTAC GCCAAGGAGGGTCCACCCGAAGCTCAACCAGATGGACCAGGACAAGGTGGCCAAGATGTACAGTGACCTGAGG AAAGAATCTATGGCGACAGGCAGCATCCCCATTACGGTGCGGCACATCGAGTCCATGAGTCATGGCGGAGGCCCA CGCGCGCATCCATCTGCGGGACTATGTGATCGAAGACGACGTCAACATGGCCATCCGCGTGATGCTGGAGAGCTT AACAATGAGCTGTTGCTCTTCATACTGAAGCAGTTAGTGGCAGAGCAGGTGACATATCAGCGCAACCGCTTTGGG GCCCAGCAGGACACTATTGAGGTCCCTGAGAAGGACTTGGTGGATAAGGCTCGTCAGATCAACATCCACAACCTC TCTGCATTTTATGACAGTGAGCTCTTCAGGATGAACAAGTTCAGCCACGACCTGAAAAGGAAAATGATCCTGCAG  $\texttt{CAGTTC} \underline{\textbf{TGA}} \texttt{GGCCCTATGCCATAAGGATTCCTTGGGATTCTGGTTTTGGGGTGGTCAGTGCCCTCTGTGCTT}$ TATGGACACAAAACCAGAGCACTTGATGAACTCGGGGTACTAGGGTCAGGGCTTATAGCAGGATGTCTGGCTGCA TTACTTGGTTGCTGAACATCTTGCCACCTCCGAGTGCTTTGTCTCCACTCAGTACCTTGGATCAGAGCTGCTGAG TTCAGGATGCCTGCGTGTGTTTAGGTGTTAGCCTTCTTACATGGATGTCAGGAGAGCTGCTGCCCTCTTGGCGT GAGTTGCGTATTCAGGCTGCTTTTGCTCGCTTTGGCCAGAGAGCTGGTTGAAGATGTTTGTAATCGTTTTCAGTC  ${\tt TCCTGCAGGTTTCTGTGCCCCTGTGGTGGAAGAGGCACGACAGTGCCAGCGCAGCGTTCTGGGCTCCTCAGTCGC}$ AAAA

#### 499/2825 FIGURE 452

MASSPAQRRRGNDPLTSSPGRSSRRTDALTSSPGRDLPPFEDESEGLLGTEGPLEEEEDGEELIGDGMERDYRAI
PELDAYEAEGLALDDEDVEELTASRREAADGPCGTVTGSWPGLGACAVGSCMTAMRRTRSALPASAASGAGTEDG
EEDEQMIESIENLEDLKGHSVREWVSMAGPRLEIHHRFKNFLRTHVDSHGHNVFKERISDMCKENRESLVVNYED
LAAREHVLAYFLPEAPAELLQIFDEAALEVVLAMYPKYDRITNHIHVRISHLPLVEELRSLRQLHLNQLIRTSGV
VTSCTGVLPQLSMVKYNCNKCNFVLGPFCQSQNQEVKPGSCPECQSAGPFEVNMEETIYQNYQRIRIQESPGKVA
ARRLPRSKDAILLADLVDSCNAGDEIELTGIYHNNYDGSLNTANGFPVFATVILANHVAKKDNKVAVGELTDEDV
KMITSLSKDQQIGEKIFASIAPSIYGHEDIKRGPALALFGGEPKNPGGKHKVRGDINVLLCGDPGTAKSQFLKYI
EKVSSRAIFTTGQGASAVAVTAYVQRHPVSREWTLEAGALVLADRGVCLIDEFDKMNDQDRTSIHEAMEQQSISI
SKAGIVTSLQARCTVIAAANPIGGRYDPSLTFSENVDLTEPIISRFDILCVVRDTVDPVQDEMLARFVVGSHVRH
HPSNKEEEGLANGSAAEPAMPNTYGVEPLPQEVLKKYIIYAKERVHPKLNQMDQDKVAKMYSDLRKESMATGSIP
ITVRHIESMSHGGGPRAHPSAGLCDRRRRQHGHPRDAGELHRHTEVQRHRSMRKTFARYLSFRRDNNELLLFILK
QLVAEQVTYQRNRFGAQQDTIEVPEKDLVDKARQINIHNLSAFYDSELFRMNKFSHDLKRKMILQQF

#### 500/2825 FIGURE 453

 $\tt CCGGTGCCGGGCGAAC{\color{red}{\textbf{ATG}}} GCGCGCCCCCGGACCCTCGTTTTGGCTGGGGAATGAAACCCTGAAGGTGCCGCT$ GGCGCTCTTTGCCTTGAACCGGCAGCGCCTGTGTGAGCGGCTGCGGAAGAACCCTGCTGTGCAGGCCGGCTCCAT CGTGGTCCTGCAGGGCGGGAGGAGACTCAGCGCTACTGCACCGACACCGGGGTCCTCTTCCTCCAGGAGTCCTT CTTTCACTGGGCGTTCGGTGTCACTGAGCCAGGCTGCTATGGTGTCATCGATGTTGACACTGGGAAGTCGACCCT GTTTGTGCCCAGGCTTCCTGCCAGCCATGCCACCTGGATGGGAAAGATCCATTCCAAGGAGCACTTCAAGGAGAA GTATGCCGTGGACGACGTCCAGTACGTAGATGAGATTGCCAGCGTCCTGACGTCACAGAAGCCCTCTGTCCTC CACTTTGCGTGGCGTCAACACGGACAGCGGCAGTGTCTGCAGGGAGGCCTCCTTTGACGGCATCAGCAAGTTCGA AGTCAACAATACCATTCTTCACCCAGAGATCGTTGAGAGCCGAGTGTTTAAGACGGATATGGAGCTGGAGGTTCT GCGCTATACCAATAAAATCTCCAGCGAGGCCCACCGTGAGGTAATGAAGGCTGTAAAAGTGGGAATGAAAGAATA TGGGTTGGAAAGCCTCTTCGAGCACTACTGCTACTCCCGGGGCGGCATGCGCCACAGCTCCTACACCTGCATCTG CGGCAGTGGTGAGAACTCAGCCGTGCTACACTACGGACACGCCGGAGCTCCCAACGACCGAACGATCCAGAATGG GGATATGTGCCTGTTCGACATGGGCGGTGAGTATTACTCTGTCGCTTCCGACATCACCTGCTCCTTTCCCCGCAA CGGCAAGTTCACTGCAGACCAGAAGGCCGTCTATGAGGCAGTGCTGCTGAGCTCCCGTGCCGTCATGGGTGCCAT GAAGCCAGGTGACTGGTGGCCTGACATCGACCGCCTGGCTGACCGCATCCACCTGGAGGAGCTGGCCCACATGGG CATCCTGAGCGCCAGCGTGGACGCCATGGTCCAGGCTCACCTGGGGGCCGTGTTTATGCCTCACGGGCTTGGCCA CTTCCTGGGCATTGACGTGCACGTGGGAGGCTACCCAGAGGGCGTGGAGCGCATCGACGAGCCCGGCCTGCG GAGCCTGCGCACTGCACGCCAGGCATGGTGCTCACCGTGGAGCCGGGCATCTACTTCATCGACCA TTTTGGCGGGGTCCGCATCGAGGAGGACGTCGTGGTGATCGACAGCGGCATAGAGCTGCTGACCTGCGTGCCCCG CACTGTGGAAGAGTTGAAGCATGCATGGCAGGCTGTGACAAGGCCTTTACCCCCTTCTCTGGCCCCAAG**TAG**AG CCAGCCAGAAATCCCAGCGCACCTGGGGGCCTGGCCTTGCAACCTCTTTTCGTGATGGGCAGCCTGCTGGTCAGC ACTCCAGTAGCGAGAGACGGCACCCAGAATCAGATCCCAGCTTCGGCATTTGATCAGACCAAACAGTGCTGTTTC  ${\tt CCGGGGAGGAAACACTTTTTAATTACCCTTTTGCAGGCACCACCTTTAATCTGTTTTATACCTTGCTTATTAAA}$ TGAGCGACTTAAAATGATTGAAAATAATGCTGTCCTTTAGTAGCAAGTAAAATGTGTCTTGCTGTCATTTATATT GTTTCCTAAAATG

## 501/2825 FIGURE 454

MAAATGPSFWLGNETLKVPLALFALNRQRLCERLRKNPAVQAGSIVVLQGGEETQRYCTDTGVLFLQESFFHWAF
GVTEPGCYGVIDVDTGKSTLFVPRLPASHATWMGKIHSKEHFKEKYAVDDVQYVDEIASVLTSQKPSVLLTLRGV
NTDSGSVCREASFDGISKFEVNNTILHPEIVESRVFKTDMELEVLRYTNKISSEAHREVMKAVKVGMKEYGLESL
FEHYCYSRGGMRHSSYTCICGSGENSAVLHYGHAGAPNDRTIQNGDMCLFDMGGEYYSVASDITCSFPRNGKFTA
DQKAVYEAVLLSSRAVMGAMKPGDWWPDIDRLADRIHLEELAHMGILSGSVDAMVQAHLGAVFMPHGLGHFLGID
VHDVGGYPEGVERIDEPGLRSLRTARHLQPGMVLTVEPGIYFIDHLLDEALADPARASFLNREVLQRFRGFGGVR
IEEDVVVIDSGIELLTCVPRTVEEIEACMAGCDKAFTPFSGPK

#### 502/2825 FIGURE 455A

GTGATAACCCCCGTGAGGACAGGCCATGGCTACGTATACGAGTACCCATCCAGATACCAAAAGGACGTCTATGAT CCAGTGGGAGAGATAAAACCTCAAGGGGTGTATGACATCCCGCCTACAAAAGGGGGTATATGCCATTCCGCCCTCT GCTTGCCGGGATGAAGCAGGGCTTAGGGAAAAAGACTATGACTTCCCCCCTCCCATGAGACAAGCTGGAAGGCCG GACCTCAGACCGGAGGGGGTTTATGACATTCCTCCAACCTGCACCAGGCAGCAGGGAAGGACCTTCATGTAAAA CCGCAACTCGGACAGTCAGTGGGCTCTCAGAACGACGCATATGATGTCCCCCGAGGCGTTCAGTTTCTTGAGCCA CCAGCAGAAACCAGTGAGAAAGCAAACCCCCAGGAAAGGGATGGTGTTTATGATGTCCCTCTGCATAACCCGCCA GATGCTAAAGGCTCTCGGGACTTGGTGGATGGGATCAACCGATTGTCTTTCTCCAGTACAGGCAGCACCCGGAGT AACATGTCCACGTCTTCCACCTCCTCCAAGGAGTCCTCACTGTCAGCCTCCCCAGCTCAGGACAAAAGGCTCTTC CTGGATCCAGACACAGCTATTGAGAGACTTCAGCGGCTCCAGCAGGCCCTTGAGATGGGTGTCTCCAGCCTAATG GCACTGGTCACTACCGACTGGCGGTGTTACGGATATATGGAAAGACACATCAATGAAATACGCACAGCAGTGGAC CTCATCCTCCACAACAAGATGAAGCGGGAGCTGCAACGAGTTGAAGACTCCCACCAGATCCTGAGTCAAACCAGC CATGACTTAAATGAGTGCAGCTGGTCCCTGAATATCTTGGCCATCAACAAGCCCCAGAACAAGTGTGACGATCTG GAGGCCCTCTTCAGACCCGGCCCTGGCAGCTTGCATCTGAAGAATGGGCCGGAGAGCATCATGAACTCAACGGAG TACCCACACGGTGGCTCCCAGGGACAGCTGCTGCATCCTGGTGACCACAAGGCCCAGGCCCCACAACAAGGCACTG TACGATTACGTCCACCTACAGGGTAAGGAGGGTTTGAGAGGCCAACAGAAGAGCTATTGGAAAAAGAGAATATC ATGAAACAGAACAAGATGCAGCTGGAACATCATCAGCTGAGCCAGTTCCAGCTGTTGGAACAAGAGATTACAAAG CCCGTGGAGATGACATCTCGAAGTGGAAGCCCTCTCAGAGCCTACCCACCACAAACAGTGGCGTGAGTGCTCAG GATCGCCAGTTGCTGTGCTTCTACTATGACCAATGTGAGACCCATTTCATTTCCCTTCTCAACGCCATTGACGCA CTCTTCAGTTGTCAGCTCAGCCCAGCCCCGCGAATCTTCGTGGCACACAGCAAGTTTGTCATCCTCAGTGCA CACAAACTGGTGTTCATTGGAGACACGCTGACACGGCAGGTGACTGCCCAGGACATTCGCAACAAGTCATGAAC TCCAGCAACCAGCTCTGCGAGCAGCTCAAGACCATAGTCATGGCAACCAAGATGGCCGCCCTCCATTACCCCAGC ACCACGGCCCTGCAGGAAATGGTGCACCAAGTGACAGACCTTTCTAGAAATGCCCAGCTGTTCAAGCGCTCTTTG CTGGAGATGGCAACGTTC**TGA**GAAGAAAAAAAAGAGGAAGGGGGCTTAACGGTTACTAAGGAAAACTGGAA ATACTGTCTGGTTTTTGTAAATGTTATCTATTTTTGTAGATATTTTATATAAAAATGAAATATTTTAACATTTTA TATACATAGAAGTATCTGAGACATAAACTGTACAGAAAACTTGTCCACGTGCTTTTGTATGCCCATGTATTCATG TTCTAATGCTGCATTTTTTTTTTTTTTTTTTAAAAACATACCAGCTGGTTATAATATTGTTCTCCACGTCCTTGTG ATGATTCTGAGCCTGGCACTCCCAAATCTGGGAAGCATAGTTTATTTGCAAGTGTTCACCTTCCAAATCATGAGG CATAGCATGACTTATTCTTGTTTGGAAAACTCTTTTCAAAACTGACCATCTTAAACACATGATGGCCAAGTGCCC AAAAGCCCTCTTGCGGAGCAAATTTCAGAATATATATGTGGATCCAAGCTCTGATAGTTCAGGTGCTGGAGGGAA GAGAGACCTGTGTGTTTAGAGGCCAGGACCACAGTTAGGATTGGGTTGTTTCAATACTGAGAGACAGCTACAATA AAAGGAGACAATTGCCTCCCTGGGGCTGTTCAATCTTCTGCATTTGTGAGTGGTTCAGTCATGAGGTTTTCCAA AAGATGTTTTTAGAGTTGTAAAAACCATATTTGCAGCAAAGATTTACAAAGGCGTATCAGACTATGATTGTTCAC CAAAATAGGGGAATGGTTTGATCCGCCAGTTGCAAGTAGAGGCCTTTCTGACTCTTAATATTCACTTTGGTGCTA CTACCCCCATTACCTGAGGGAAACTGGCCAGGTCCTTGATCATGGAACTATAGAGCTACCAGGACATATCCTGCT  $\tt CTCTAAGGGAATTTATTGCTATCTTGCACCTTCTTTAAAACTCACATATGCAGACCTGACACTCAAGAGTGGCTA$ GCTACACAGAGTCCATCTAATTTTTGCAACTTCCTGTGGCCAGTGTGTATAACCCCTTCCACTATCTCACAGATA GTCACAGCGTCCATTCCATAGTCTGTCTCCTCACATCTGTTAGTATTGACACAGCACAGACACCACAAGCCATCA GGTTCTTCATGGGGCAGGTGAAATACTTCTACCCCATGGGTAAATGTATTCACATATTACCAAGAAGAAGAACAC ATTATCTATGATCTTTTGGCCCAGTTCTTATTTAGCATTTTTATTCCAGCCTACTTGGAAACATGTTTTTATTTG CAATATATGCCTGACTGAATTAAGCTTGCTTGTTTTAAACAACCAAATCATTGGAACAGAAAAGGATTTAAAAAA 

# 503/2825 FIGURE 455B

# 504/2825 FIGURE 456

VITPVRTGHGYVYEYPSRYQKDVYDIPPSHTTQGVYDIPPSSAKGPVFSVPVGEIKPQGVYDIPPTKGVYAIPPS
ACRDEAGLREKDYDFPPPMRQAGRPDLRPEGVYDIPPTCTKPAGKDLHVKYNCDIPGAAEPVARRHQSLSPNHPP
PQLGQSVGSQNDAYDVPRGVQFLEPPAETSEKANPQERDGVYDVPLHNPPDAKGSRDLVDGINRLSFSSTGSTRS
NMSTSSTSSKESSLSASPAQDKRLFLDPDTAIERLQRLQQALEMGVSSLMALVTTDWRCYGYMERHINEIRTAVD
KVELFLKEYLHFVKGAVANAACLPELILHNKMKRELQRVEDSHQILSQTSHDLNECSWSLNILAINKPQNKCDDL
DRFVMVAKTVPDDAKQLTTTINTNAEALFRPGPGSLHLKNGPESIMNSTEYPHGGSQGQLLHPGDHKAQAHNKAL
PPGLSKEQAPDCSSSDGSERSWMDDYDYVHLQGKEEFERQQKELLEKENIMKQNKMQLEHHQLSQFQLLEQEITK
PVENDISKWKPSQSLPTTNSGVSAQDRQLLCFYYDQCETHFISLLNAIDALFSCVSSAQPPRIFVAHSKFVILSA
HKLVFIGDTLTRQVTAQDIRNKVMNSSNQLCEQLKTIVMATKMAALHYPSTTALQEMVHQVTDLSRNAQLFKRSL
LEMATF

#### 505/2825 FIGURE 457

AGATCGGGGCCAAGTTTTGGGAGGTTATCAGTGACGAACATGGCATCGACCCCACAGGCACATACCATGGGGACA GTGACCTGCAACTGGAGAGGATCAACGTGTACTACAACGAGGCCACAGGAGGAAATTATGTCCCCAGAGCGGTGC TGGTGGACCTGGAACCCGGCACCATGGACTCTGTCCGTTCTGGCCCCTTCGGTCAGATCTTTCGGCCGGACAACT TCGTGTTTGGCCAATCCGGAGCCGGCAACAACTGGGCAAAGGGGCACTACACGGAGGGCGCAGAGCTGGTGGACG CTGTCCTGGACGTAGTCCGGAAGGAGGCCGAGAGCTGCGACTGCCTTCAGGGGCTTCCAGCTGACCCACTCGCTGG GGGGTGGCACGGGGTCCGGAATGGGCACGCTGCTCATCAGTAAGATCCGCGAGGAGTTCCCAGACCGCATCATGA ACCAGCTGGTGGAGAATACGGATGAGACCTACTGCATCGACAACGAGGCACTCTACGACATCTGTTTCCGCACCC TCAAGCTGACCACCCCACCTACGGGGACCTCAACCACCTGGTGTCGGCCACCATGAGCGGGGTCACCACCTGCC TGCGCTTCCCGGGCCAGCTGAACGCCGACCTGCGCAAGCTGGCCGTCAACATGGTTCCCTTTCCTCGCCTGCACT CCCAGCAGATGTTCGATGCCAAGAACATGATGGCGGCGTGCGACCCGCCCACGGCCGCTACCTGACCGTGGCCG CCGTGTTCCGGGGCCGCATGTCCATGAAGGAGGTGGACGAGCAGATGCTGAGCGTGCAGAGCAAGAACAGCAGCT ACTTCGTGGAGTGGATCCCCAACAACGTGAAGACGGCCGTGTGCGACATCCCGCCCCGCGGCCTGAAGATGGCCG CGACCTTCATCGGCAACAGCACGGCCATCCAGGAGCTGTTCAAGCGCATCTCCGAGCAGTTCACGGCCATGTTCC GGCGCAAGGCCTTCTTGCACTGGTACACGGGCGAGGGCATGGACGAGTTGACTTCACCGAGGCCGAGAGCAACA TGAATGACCTGGTATCTGAGTACCAGCAGTACCAGGACGCCACGGCCGAGGAGGGCGAGTTCGAGGAGGAGGCGG AGGAGGAGGTGGCC**TAG**GCTGCTCCCATCGCTTCCCACCTGTCCCCTCGAGGCTTCTGACCTTTGATCCGCTAGG CCCCCATCTCTGAACCCTAGAGCCCCGCTTTCCCTCCAAGGCTGACTCCCCGCTGACCCTAACAATACCTTTGG AGCTCGCTTTACCTCTGGCTACTTCATCTCCGACCCTGGCTCCCCTTTGAGCCCTAATTTATCTTTAACCCCCTT GAGCTCTTCCAACCTTGACATTCCCAGGAGGAGCCCCGCTTCACCCCTTCTGACTCTGGAAACCGCACCTTTAAC TTTGCAGACCTTCCTTCACCCCTGACTTCTGCTTCACCTTTGACCTCTGCCCCCCATGAATCCCATTTTACCTCT AGACCTATAAGTTCTGGTTTATGTTTGACCCCTCCCTCTGAGCTGCACTTCACCGCTGACCTTGCCTCACCTTTA ACCCCCACCTGAGCCCCAGCTCCTACCTCTGACCCCAACTTCTCTTTGATCTCTGAATCCCCTCTGACTCCAAC CCAGCTTCTCTGATTCCCACAGGCCCCATGCATCCTCCCTGCCTCACTCCCTCAGCCCCTGCCGACCTTAGC 

## 506/2825 FIGURE 458

MREIVHLQAGQCGNQIGAKFWEVISDEHGIDPTGTYHGDSDLQLERINVYYNEATGGNYVPRAVLVDLEPGTMDS VRSGPFGQIFRPDNFVFGQSGAGNNWAKGHYTEGAELVDAVLDVVRKEAESCDCLQGFQLTHSLGGGTGSGMGTL LISKIREEFPDRIMNTFSVVPSPKVSDTVVEPYNATLSVHQLVENTDETYCIDNEALYDICFRTLKLTTPTYGDL NHLVSATMSGVTTCLRFPGQLNADLRKLAVNMVPFPRLHFFMPGFAPLTSRGSQQYRALTVPELTQQMFDAKNMM AACDPRHGRYLTVAAVFRGRMSMKEVDEQMLSVQSKNSSYFVEWIPNNVKTAVCDIPPRGLKMAATFIGNSTAIQ ELFKRISEQFTAMFRRKAFLHWYTGEGMDEMEFTEAESNMNDLVSEYQQYQDATAEEGEFEEEAEEEVA

### 507/2825 FIGURE 459A

TGAGGAATCAACAGCCGCCATCTTGTCGCGGACCCGACCGGGGCTTCGAGCGCGATCTACTCGGCCCCGCCGGTC CCGGGCCCCACAACCGCCCGCGCTCGCTCTCCCTCGCAGCCGGCAGGGCCCCCGACCCCGTCCGGGCCCTC CCCAACCCCAAAAGAGCCAAACTCAGCTCGCCCGGTTTCTCGGCGAATGACAGCACAGATTTTGGATCATTGTTT GACTTGGAAAATGATCTTCCTGATGAGCTGATACCCAATGGAGGAGAATTAGGCCTTTTAAACAGTGGGAACCTT GTTCCAGATGCTGCTTCCAAACATAAACAACTGTCGGAGCTTCTACGAGGAGGCAGCGGCTCTAGTATCAACCCA GGAATAGGAAATGTGAGCGCCAGCAGCCCCGTGCAGCAGGGCCTGGGTGGCCAGGCTCAAGGGCAGCCGAACAGT AAACAGGCAGCCAGCCTCTGGGCCCACCCCGCTGCCTCCCAAGCACTGAATCCGCAAGCACAAAAGCAAGTG GGGCTGGCGACTAGCAGCCCTGCCACGTCACAGACTGGACCTGGTATCTGCATGAATGCTAACTTTAACCAGACC AATGGATCTCTTGGGGCTGCTGGCAGAGGAAGGGGAGCTGGAATGCCGTACCCTACTCCAGCCATGCAGGGCGCC TCGAGCAGCGTGCTGAGACCCTAACGCAGGTTTCCCCGCAAATGACTGGTCACGCGGGACTGAACACCGCA CAGGCAGGAGGCATGGCCAAGATGGGAATAACTGGGAACACAAGTCCATTTGGACAGCCCTTTAGTCAAGCTGGA GGGCAGCCAATGGGAGCCACTGGAGTGAACCCCCAGTTAGCCAGCAAACAGAGCATGGTCAACAGTTTGCCCACC TTCCCTACAGATATCAAGAATACTTCAGTCACCAACGTGCCAAATATGTCTCAGATGCAAACATCAGTGGGAATT GTACCCACACAAGCAATTGCAACAGGCCCCACTGCAGATCCTGAAAAACGCAAACTGATACAGCAGCAGCTGGTT CTACTGCTTCATGCTCATAAGTGTCAGAGACGAGAGCAAACGGAGAGGTTCGGGCCTGCTCCCGCAT TGTCGAACCATGAAAAACGTTTTGAATCACATGACGCATTGTCAGGCTGGGAAAGCCTGCCAAGTTGCCCATTGT GCATCTTCACGACAAATCATCTCTCATTGGAAGAACTGCACACGACATGACTGTCCTGTTTGCCTCCCTTTGAAA AATGCCAGTGACAAGCGAAACCAACAAACCATCCTGGGGTCTCCAGCTAGTGGAATTCAAAACACAATTGGTTCT GTTGGCACAGGGCAACAGAATGCCACTTCTTTAAGTAACCCAAATCCCATAGACCCCAGCTCCATGCAGCGAGCC TATGCTGCTCTCGGACTCCCCTACATGAACCAGCCCCAGACGCAGCTGCAGCCTCAGGTTCCTGGCCAGCAACCA GCACAGCCTCAAACCCACCAGCAGATGAGGACTCTCAACCCCCTGGGAAATAATCCAATGAACATTCCAGCAGGA GGAATAACAACAGATCAGCAGCCCCCAAACTTGATTTCAGAATCAGCTCTTCCGACTTCCCTGGGGGCCCACAAAC CCACTGATGAACGATGGCTCCAACTCTGGTAACATTGGAACCCTCAGCACTATACCAACAGCAGCTCCTCCTTCT AGCACCGGTGTAAGGAAAGGCTGGCACGAACATGTCACTCAGGACCTGCGGAGCCATCTAGTGCATAAACTCGTC CAAGCCATCTTCCCAACACCTGATCCCGCAGCTCTAAAGGATCGCCGCATGGAAAACCTGGTAGCCTATGCTAAG AAAGTGGAAGGGGACATGTACGAGTCTGCCAACAGCAGGGATGAATATTATCACTTATTAGCAGAGAAAATCTAC TTACCAGCCCCGGGGGCTCAGCCCCTGTGATTCCACAGGCACAACCTGTGAGACCTCCAAATGGACCCCTGTCC CTGCCAGTGAATCGCATGCAAGTTTCTCAAGGGATGAATTCATTTAACCCCATGTCCTTGGGGAACGTCCAGTTG CCACAAGCACCCATGGGACCTCGTGCAGCCTCCCCAATGAACCACTCTGTCCAGATGAACAGCATGGGCTCAGTG GCCCAGGCGCCCGCTCAGAGCCAGTTTCTGCCACAGAACCAGTTCCCGTCATCCAGCGGGGCGATGAGTGTGGGC ATGGGGCAGCCGAACAGGCGTGTCACAGGGACAGGTGCCTGGTGCTCTTCCTAACCCTCTCAAC ATGCTGGGGCCTCAGGCCAGCTACCTTGCCCTCCAGTGACACAGTCACCACTGCACCCAACACCGCCTCCT CCCACTCAGCCATCAACTCCTGTGTCGTCTTCCGGGCAGACTCCCACCCCGACTCCTGGCTCAGTGCCCAGTGCT ACCCAAACCCAGAGCACCCCTACAGTCCAGGCAGCCCAGGCCCAGGTGACCCCGCAGCCTCAAACCCCAGTT CAGCCCCGTCTGTGGCTACCCCTCAGTCATCGCAGCAACAGCCGACGCCTGTGCACGCCCAGCCTCCTGGCACA CCGCTTTCCCAGGCAGCAGCATTGATAACAGAGTCCCTACCCCTCCTCGGTGGCCAGCAGCAGAAACCAAT TCCCAGCAGCCAGGACCTGACGTACCTGTGCTGGAAATGAAGACGGAGACCCAAGCAGAGGACACTGAGCCCGAT CCTGGTGAATCCAAAGGGGAGCCCAGGTCTGAGATGATGGAGGAGGATTTGCAAGGAGCTTCCCAAGTTAAAGAA GAAACAGACATAGCAGAGCAGAAATCAGAACCAATGGAAGTGGATGAAAAGAAACCTGAAGTGAAAGTAGAAGTT AAAGAGGAAGAAGAGTAGCAGTAACGGCACAGCCTCTCAGTCAACATCTCCTTCGCAGCCGCGCAAAAAAATC TTTAAACCAGAGGAGTTACGCCAGGCCCTCATGCCAACCCTAGAAGCACTGTATCGACAGGACCCAGAGTCATTA CCTTTCCGGCAGCCTGTAGATCCCCAGCTCCTCGGAATTCCAGACTATTTTGACATCGTAAAGAATCCCATGGAC

#### 508/2825 FIGURE 459B

 $\verb|CTCTCCACCATCAAGCGGAAGCTGGACACAGGGCAATACCAAGAGCCCTGGCAGTACGTGGACGACGTCTGGCTC| \\$ ATGTTCAACAATGCCTGGCTCTATAATCGCAAGACATCCCGAGTCTATAAGTTTTGCAGTAAGCTTGCAGAGGTC ACTTTGTGCTGCTATGGGAAGCAGCTGTTGTACCATTCCTCGCGATGCTGCCTACTACAGCTATCAGAATAGGTAT ACGACAATTTCAAAGGATCAGTTTGAAAAGAAGAAAAATGATACCTTAGACCCCGAACCTTTCGTTGATTGCAAG GAGTGTGGCCGGAAGATGCATCAGATTTGCGTTCTGCACTATGACATCATTTGGCCTTCAGGTTTTGTGTGCGAC AACTGCTTGAAGAAACTGGCAGACCTCGAAAAGAAAACAAATTCAGTGCTAAGAGGCTGCAGACCACAAGACTG GGAAACCACTTGGAAGACCGAGTGAACAAATTTTTGCGGCGCCAGAATCACCCTGAAGCCGGGGAGGTTTTTGTC CGAGTGGTGGCCAGCTCAGACAAGACGGTGGAGGTCAAGCCCGGGATGAAGTCACGGTTTGTGGATTCTGGGGAA ATGTCTGAATCTTTCCCATATCGAACCAAAGCTCTGTTTGCTTTTGAGGAAATTGACGGCGTGGATGTCTGCTTT TTTGGAATGCACGTCCAAGAATACGGCTCTGATTGCCCCCCTCCAAACACGAGGCGTGTGTACATTTCTTATCTG GATAGTATTCATTTCTTCCGGCCACGTTGCCTCCGCACAGCCGTTTACCATGAGATCCTTATTGGATATTTAGAG TATGTGAAGAAATTAGGGTATGTGACAGGGCACATCTGGGCCTGTCCTCCAAGTGAAGGAGATGATTACATCTTC CATTGCCACCCACCTGATCAAAAAATACCCAAGCCAAAACGACTGCAGGAGTGGTACAAAAAGATGCTGGACAAG GCGTTTGCAGAGCGGATCATCCATGACTACAAGGATATTTTCAAACAAGCAACTGAAGACAGGCTCACCAGTGCC AAGGAACTGCCCTATTTTGAAGGTGATTTCTGGCCCAATGTGTTAGAAGAGAGCATTAAGGAACTAGAACAAGAA GAAGAGGAGAAAAAGGAAGAGACACTGCAGCCAGTGAAACCACTGAGGGCAGTCAGGGCGACAGCAAGAAT ATGCCCAACGTGTCCAATGACCTGTCCCAGAAGCTGTATGCCACCATGGAGAAGCACAAGGAGGTCTTCTTCGTG ATCCACCTGCACGCTGGGCCTGTCATCAACACCCTGCCCCCATCGTCGACCCCGACCCCTGCTCAGCTGTGAC CTCATGGATGGGCGCGACGCCTTCCTCACCCTCGCCAGAGACAAGCACTGGGAGTTCTCCTCCTTGCGCCGCTCC AAGTGGTCCACGCTCTGCATGCTGGAGCTGCACACCCAGGGCCAGGACCGCTTTGTCTACACCTGCAACGAG TGCAAGCACCACGTGGAGACGCGCTGGCACTGCACTGTGTGCGAGGACTACGACCTCTGCATCAACTGCTATAAC CAGTCAAAGAGCCCCCAGGAGTCACGCCGGGTGAGCATCCAGCGCTGCATCCAGTCGCTGGTGCACGCGTGCCAG TGCCGCAACGCCAACTGCTCGCCATCCTGCCAGAAGATGAAGCGGGTGGTGCAGCACCAAGGGCTGCAAA CGCAAGACCAACGGGGGCTGCCCGGTGTGCAAGCAGCTCATCGCCCTCTGCTGCTACCACGCCAAGCACTGCCAA GAAAACAAATGCCCCGTGCCCTTCTGCCTCAACATCAAACACAAGCTCCGCCAGCAGCAGATCCAGCACCGCCTG  ${\tt CAGCAGGCCCAGCTCATGCGCCGGATGGCCACCATGAACACCCGCAACGTGCCTCAGCAGAGTCTGCCTTCT}$ CGGCAGATCGAGCGTGAGGCCCAGCAGCAGCACCTGTACCGGGTGAACATCAACAACAGCATGCCCCCAGGA CGCACGGGCATGGGGACCCCGGGGAGCCAGATGGCCCCCGTGAGCCTGAATGTGCCCCGACCCAACCAGGTGAGC AGGAGCATCTCACCCAGCGCTCTGCAAGACCTGCTGCGGACCCTGAAGTCGCCCAGCTCCCCTCAGCAGCAACAG CAGGTGCTGAACATCTCAAATCAAACCCGCAGCTAATGGCAGCTTTCATCAAACAGCGCACAGCCAAGTACGTG GCCAATCAGCCCGGCATGCAGCCCCAGCCTGGCCTCCAGTCCCAGCCCGGCATGCAACCCCAGCCTGGCATGCAC CAGCAGCCCAGCCTGCAGAACCTGAATGCCATGCAGGCTGGCGTGCCGGGGCCCGGTGTGCCTCCACAGCAGCAG GCGATGGGAGGCCTGAACCCCCAGGGCCAGGCCTTGAACATCATGAACCCAGGACACAACCCCAACATGGCGAGT ATGAATCCACAGTACCGAGAAATGTTACGGAGGCAGCTGCTGCAGCAGCAGCAGCAGCAGCAGCAACAACAG CAGCAACAGCAGCAGCAAGGGAGTGCCGGCATGGCTGGGGGCATGGCGGGGCACGGCCAGTTCCAGCAGCCT CAAGGACCCGGAGGCTACCCACCGGCCATGCAGCAGCAGCAGCATGCAGCAGCATCTCCCCCTCCAGGGCAGC TCCATGGGCCAGATGGCGGCTCAGATGGGACAGCTTGGCCAGATGGGGCAGCCGGGGCTGGGGGCAGACAGCACC CCCAACATCCAGCAGCCCTGCAGCAGCGGATTCTGCAGCAACAGCAGATGAAGCAGCAGATTGGGTCCCCAGGC CAGCCGAACCCCATGAGCCCCCAGCAACACATGCTCTCAGGACAGCCACAGGCCTCGCATCTCCCTGGCCAGCAG

#### 509/2825 FIGURE 459C

ATCGCCACGTCCCTTAGTAACCAGGTGCGGTCTCCAGCCCCTGTCCAGTCTCCACGGCCCCAGTCCCAGCCTCCA CATTCCAGCCCGTCACCACGGATACAGCCCCAGCCTTCGCCACACCACGTCTCACCCCAGACTGGTTCCCCCCAC CCCGGACTCGCAGTCACCATGGCCAGCTCCATAGATCAGGGACACTTGGGGAACCCCGAACAGAGTGCAATGCTC  $\verb|CCCCAGCTGAACACCCCCAGCAGGAGTGCGCTGTCCAGCGAACTGTCCCTGGTCGGGGACACCACGGGGGGACACG|$ CTAGAGAAGTTTGTGGAGGGCTTGTAGCATTGTGAGAGCATCACCTTTTCCCTTTCATGTTCTTGGACCTTTTGT ACTGAAAATCCAGGCATCTAGGTTCTTTTTATTCCTAGATGGAACTGCGACTTCCGAGCCATGGAAGGGTGGATT TTGTTTGGATTCTTTTTGTCGTCATTGCTGGTGACTCATGCCTTTTTTTAACGGGAAAAACAAGTTCATTATATT GATTCCTGAAATAATGCAAGCTTATAATGTATCCCGATAACTTTGTGATGTTTCGGGAAGATTTTTTTCTATAGT GAACTCTGTGGGCGTCTCCCAGTATTACCCTGGATGATAGGAATTGACTCCGGCGTGCACACACGTACACCCCA CACACATCTATCTATACATAATGGCTGAAGCCAAACTTGTCTTGCAGATGTAGAAATTGTTGCTTTGTTTCTCTG ATAAAACTGGTTTTAGACAAAAAATAGGGATGATCACTCTTAGACCATGCTAATGTTACTAGAGAAGAAGCCTTC TTTTCTTCTTCTATGTGAAACTTGAAATGAGGAAAAGCAATTCTAGTGTAAATCATGCAAGCGCTCTAATTCCT ATAAATACGAAACTCGAGAAGATTCAATCACTGTATAGAATGGTAAAATACCAACTCATTTCTTATATCATATTG TTAAATAAACTGTGTGCAACAGACAAAAAGGGTGGTCCTTCTTGAATTCATGTACATGGTATTAACACTTAGTGT TCGGGGTTTTTTGTTATGAAAATGCTGTTTTCAACATTGTATTTGGACTATGCATGTGTTTTTTCCCCATTGTAT ATAAAGTACCGCTTAAAATTGATATAAATTACTGAGGTTTTTAACATGTATTCTGTTCTTTAAGATCCCCTGTAA 

# 510/2825 FIGURE 460

MAENLLDGPPNPKRAKLSSPGFSANDSTDFGSLFDLENDLPDELIPNGGELGLLNSGNLVPDAASKHKOLSELLR GGSGSSINPGIGNVSASSPVQQGLGGQAQGQPNSANMASLSAMGKSPLSQGDSSAPSLPKQAASTSGPTPAASQA LNPQAQKQVGLATSSPATSQTGPGICMNANFNQTHPGLLNSNSGHSLINQASQGQAQVMNGSLGAAGRGRGAGMP YPTPAMQGASSSVLAETLTQVSPQMTGHAGLNTAQAGGMAKMGITGNTSPFGQPFSQAGGQPMGATGVNPQLASK QSMVNSLPTFPTDIKNTSVTNVPNMSQMQTSVGIVPTQAIATGPTADPEKRKLIQQQLVLLLHAHKCQRREQANG EVRACSLPHCRTMKNVLNHMTHCQAGKACQVAHCASSRQIISHWKNCTRHDCPVCLPLKNASDKRNQOTILGSPA SGIQNTIGSVGTGQQNATSLSNPNPIDPSSMQRAYAALGLPYMNQPQTQLQPQVPGQQPAQPQTHQQMRTLNPLG NNPMNIPAGGITTDQQPPNLISESALPTSLGATNPLMNDGSNSGNIGTLSTIPTAAPPSSTGVRKGWHEHVTQDL RSHLVHKLVQAIFPTPDPAALKDRRMENLVAYAKKVEGDMYESANSRDEYYHLLAEKIYKIOKELEEKRRSRLHK QGILGNQPALPAPGAQPPVIPQAQPVRPPNGPLSLPVNRMQVSQGMNSFNPMSLGNVQLPQAPMGPRAASPMNHS VQMNSMGSVPGMAISPSRMPQPPNMMGAHTNNMMAQAPAQSQFLPQNQFPSSSGAMSVGMGQPPAQTGVSQGQVP GAALPNPLNMLGPQASQLPCPPVTQSPLHPTPPPASTAAGMPSLQHTTPPGMTPPQPAAPTQPSTPVSSSGQTPT PTPGSVPSATQTQSTPTVQAAAQAQVTPQPQTPVQPPSVATPQSSQQQPTPVHAQPPGTPLSQAAASIDNRVPTP SSVASAETNSQQPGPDVPVLEMKTETQAEDTEPDPGESKGEPRSEMMEEDLQGASQVKEETDIAEQKSEPMEVDE KKPEVKVEVKEEEESSSNGTASQSTSPSQPRKKIFKPEELRQALMPTLEALYRQDPESLPFRQPVDPQLLGIPDY FDIVKNPMDLSTIKRKLDTGQYQEPWQYVDDVWLMFNNAWLYNRKTSRVYKFCSKLAEVFEQEIDPVMQSLGYCC GRKYEFSPQTLCCYGKQLCTIPRDAAYYSYQNRYHFCEKCFTEIQGENVTLGDDPSQPQTTISKDQFEKKKNDTL DPEPFVDCKECGRKMHQICVLHYDIIWPSGFVCDNCLKKTGRPRKENKFSAKRLQTTRLGNHLEDRVNKFLRRQN HPEAGEVFVRVVASSDKTVEVKPGMKSRFVDSGEMSESFPYRTKALFAFEEIDGVDVCFFGMHVOEYGSDCPPPN TRRVYISYLDSIHFFRPRCLRTAVYHEILIGYLEYVKKLGYVTGHIWACPPSEGDDYIFHCHPPDOKIPKPKRLO EWYKKMLDKAFAERIIHDYKDIFKQATEDRLTSAKELPYFEGDFWPNVLEESIKELEQEEEERKKEESTAASETT EGSQGDSKNAKKKNNKKTNKNKSSISRANKKKPSMPNVSNDLSQKLYATMEKHKEVFFVIHLHAGPVINTLPPIV DPDPLLSCDLMDGRDAFLTLARDKHWEFSSLRRSKWSTLCMLVELHTOGODRFVYTCNECKHHVETRWHCTVCED YDLCINCYNTKSHAHKMVKWGLGLDDEGSSQGEPQSKSPQESRRVSIQRCIQSLVHACQCRNANCSLPSCQKMKR VVQHTKGCKRKTNGGCPVCKQLIALCCYHAKHCQENKCPVPFCLNIKHKLRQQQIQHRLQQAQLMRRRMATMNTR NVPQQSLPSPTSAPPGTPTQQPSTPQTPQPPAQPQPSPVSMSPAGFPSVARTQPPTTVSTGKPTSQVPAPPPPAQ PPPAAVEAARQIEREAQQQQHLYRVNINNSMPPGRTGMGTPGSQMAPVSL

### 511/2825 FIGURE 461A

GAGGATGCAGAGTGGGAGACCTTAAAACCGTCTGATCTGGCTGCTAAAGCGAGTGTCTGACGACCACCCATTAAC ACCAGCGAGTAGCGTTTGCTCACGGCTTTAGGATAAGAAGCCTGGA ATG GTCCCCCTCCGCCGGGAGAAGAAAGCCAGACGGTCATCCTTCCACCTGGCTGGCAGAGCTACCTGTCGCCTCAGGGCCGGCGGTACTATGTCAACACGAC TCTGCCTCCAACAGTGAATGGATACCACGCATCAGGGACCCCAGGCGCACCCTCCAGAGACTGCCCACATGAGTGT CCGAAAATCCACCGGTGATTCCCAGAACCTGGGATCCTCATCGCCAAGCAAAAAGCAGAGCAAGGAAAACACCAT CTACTGCGACTACTTCTGGGCTGATAAGAAGGACCCCCAAGGCAACGGCACGGTGGCTGGGTTTGAACTACTGCT CCAGAAACAGCTGAAGGGCAAACAAATGCAGAAGGAAATGTCAGAATTCATCCGGGAAAGGATAAAGATTGAAGA AGACTATGCGAAGAACTTAGCTAAGCTCTCTCAGAACTCCTTGGCTTCACAGGAGGAAGGCTCCTTGGGAGAGGC GTGGGCCCAGGTGAAGAAGAGCCTGGCGGACGAAGCAGAAGTTCACCTCAAGTTCTCTGCCAAGCTTCACAGCGA CCTTCGCAAGCAGCTCGCCAGCCGCTATGCCTCGGTGGAGAAGGCCCGGAAAGCCCTCACAGAGCGGCAGAGAGA CCTGGAGATGAAGACCCAGCAGCTGGAGATCAAGCTGAGCAACAAGACAGAGGAGGACATCAAGAAGGCGCGGAG AAAGTCCACACGGCTGGAGACGACCTCATGCGCTGTGTGGATCTCTACAACCAGGCCCAGTCCAAATGGTTTGA AGAGATGGTGACCACCACATTGGAGCTAGAGCGGCTGGAGGTGGAGAGGGTAGAGATGATCCGGCAGCACCTGTG CCAGTACACGCAGCTGCGCATGAAACAGACATGTTCAACCAAAGCACAGTCGAGCCCGTGGATCAGCTGCTTCG AAAAGTGGACCCGGCCAAAGACAGGGAGCTGTGGGTCAGAGAGCACAAGACGGGCAACATCCGCCCTGTGGACAT TGCCGAGTGGATGCCCCCCCCCCTCTCTCTGGGCCACTGAGGAGAGGGGGAGAGAGCTGGTGATTCCAGAAGGGT CTTGCCGAGTTTTGTAGGGGTCACTGTCACATGCTTGTGCCCTGGAAGCCCCAAGGCTCTTCCTCCAGCTGGCTC  ${\tt CCTTGTCTCCAGGGCTTTGGAGGATCAGGGTAGGGAGGGCTCTGTCTCTAAGCCAGGTGTCAGGATCAGAATCAT}$ GGGTAGAAGGTGCCATTCAGCTCACAGCCGCACCCAGAATCCTTTGCAGCCCTCCTTCTTTATTTTTTCCCATT GCATTCTGGGAGTCCACATCTGGCTTTCTCAGCCACTGTTCATCACCAGGGGTTTTAGGAGGAAGGCTTGGCTCC TGTCTTCCCAGACCCATGCCTGGAGAGGTCAGGATGGAACTACCTCATTCGGCAAATTAGCCCCAAATTGAG GCTGCTTCCTCGGGGAGGCCCCTGCCCACCCTGCCAGCCCTTCCCAGTCTCATTAGAGGAAGCTTTCCAAAGT AGCAAGCTGGGGCCTGTCCGCCACGGTGCCCTGTGCACCTTTGGGCTGCCCTGGCCCCCACTCTCCCGGGCCGCC CACCACAGGCTCTTAATGGGGCCGGGTCAGTCCTACATGTGAGATGGGTTAGGGCAAGTCTTTGCCATCCCCAG ATGGCTCTGTCTTGTGTATGGCAGGGCTGGGACTGCTGTCCCTTGTACAGTTTTCTGTCACTGGTGGGCACT GGACAGGCATAGCAGACTCTCTTGTGGCCACACTAGGTTGTCGCCTTTCAGGCAGTGTCCCGCCACCTTTGCTTC CCGCTCCTTCTGGACATTCCAGAGCCCTGCACCAAACCCTTACTTGGTGTCTCGCACCTCCTTTCCCTCTCCAT CTTCCAATCCCTGGAAAAGTCTGGTCTGAGTGTGACTTGGGAAGCTTTCAGTGCTGCTGTTTTGGCCCAGCTCATT CACTTTGGGTTTGACCTGCACATCTGCAGTGAGGGCAGAATTCAACAAGCACAACTCACTGGTCTTTCAGTCAAC GTGCTAGAAACCGATGACTTATTAATCTCTAATTTTTTGGCGCCTCTTTCATTGAATGAGAATTGCTTTCGTATA GTTCACATTAGAAAAATGCCTAATATACTAAAGTAAACCAAACGTTGTCACTTTTCTCTTGTTCTTGAAACATTG CAACCAAAAAGGTCAGCACAAAGGCCTTCACCTACGTGAAGACCTCCTGGTAGGATCTGTCCATGGGATGGAGAA CCACTCTGTCCAGATCTGGGGTTGGGTCATGACACCAGCTACCAATTTTAGAATATTATTTCTTGGTTTCTTTAT GAAAAATGGGTGCTAGTGGTAATTGCTTTGTGGCTTAGTAAACTACTCTGTGGATGATTTCCAAACATTCAAAGC CAATAGCCTTGTTATTAACAAGATATTTTGAGTACAATATGGCTCATTGACTTTTCCATTACATCTGAGGATTCC AGAGTCCTCTGTTCATCCCTGGGATAGAGTGAGCCCTCTTGCGTCTTCCAGGGTAGCTCAGGTTGGCCCAGTTTG GGCCAGTCCATTTTTGGAGGTCACTCTTTCCCCCCTCATCCCTCACCTCCTTCTCTTTACCCCCTTCACGTACT TCCTTTCTTCCTCCTCTGTTATTCATCATCAAGCAGGTTAAAGCCATTGTGCTAAGATCTAATCTGAGGACA

#### 512/2825 FIGURE 461B

TGATACATAGGCACTCTACAGAATTTAAGTTTTCAGAGGTAAGTCTGTTGTTCTGATTAATCTGCATTCAG CCTCTATTCTCCATGTCGACTGAGAAGAGTATAATAAGAACATTGTACCCTCTCCACACTTAACCCTCTGGGATT CACCTGGTCACCAGACCCTCACGGCCCATATCCAATTCGAATTTCAGAAATTTCTCTGTTTGCTTTAAATAACTC TTTACAAATAAGGGTCGGGCACGGTGGCTCACGCCTGTAATCCCAGCTCTTTGGGAGGCCGAGGTGGGCAGATCA CGAGGTCAGGAGATCGAGACCATCCTGGCTAACATGGTGAAACCCCGTCTCTGCTAAAAATACAAAAAATTAGCC AGATGTGGTGGTGCACACCTGTAGTCCCAACTATTCGGCAGGCTGAGGCAGTAGAATCGCTTGAACCTGGGAGGC GGAGATTGCAGTGAACCAAGATCGGGCCACTGCACTCCAGCTTGGGTGACAGAGCAAGACTCCATCTCAAAAAAA TTATAAATAACTCTTTGCAAATAGGAAAGTGAAGCTCAAGGGTAACCAGTGAGGCTATGAAATTGCCAGTGAATT AAGACCAGGGCTAGAGTCAACAGTCCAGAACCCCTATCCGTATGAACAGGGGCCCGTTTCTGGTTGAAATCAAGG ATGTCTTGGTGCTTAGGTGAGGACTTAGCTGGGTTTATTCAAAGTGGGCGGGTCTGGGCTATTCTTGAGAATGTC CCTTCCATTTCTGAGAACAAGTCAGCCCCTTCTACCTGCCTCCACCCAGGAGACTGATGCACAAGATCTGTTTGT TGGGGATGGCGGCCCTGAGATTCTGTCAGTGGGACCACCCTGGGCCTCCTTTCTACCTCCACTCAGCACCCTGTT GGCCAAGGAGAATTTCTGCGGTGGGAGGCAGTGCTCTGCTAGTCAGGATTGATAAACAGCTGGGCACACCGAAAC GGATGGATGCAGAAAATCTGGGCTGAGCCAAAGTCCCTTTTGGAAATACAAGCCATAACATTCGAAGGACATCA GCGACCTTGGCTTGTTTAGGTGATTTTACTTCCAGCTGCAGGTAGTCTTGACAAGGAGTGTTTAAACAGAAGGCT CAAGATGCATTCCTTGTGTAGGTCGGAGAGAGCACTTCTAATGTTAAGTGGGGTACAGATCAGCTGCCCCCCAC GTAGCCTGGACATCGTCTTATCCCCATAATCCTTGCCATCCCTACAAGGCCCATCGCCACCACCTTTTCCCAGGT TTATTGAGGTATGACTCAGTAAAATTCACCCTTTGGAAATATACAGCTCTGTGAATTTTGACAAATTTA GTGTCTTGTGACCATCACCAAGATCAACCTGTTTTTAACCCTCCAAAAAATTCCCTTCTCCTGCCCTCTTTCCCT GGCAACCCTGATTGATTATCTGATCCTATAATTTTGCCTTTTCTAGAATGTCATATAAATGGAGTCACACTGATG CCTTCTCCTTGCCGGGAAGTATTTCATTGCATGGGCGTACTACTCTGCTTCTTTATTTTTACACTGAGCCTTTCG CAGGCTCCTCCAAATGCCAGCTGTGCCCACAAGCTGTTCTTAGGGCTCACACTCGCCTTTTGGCGCGTGGTG TGCTGCAACCTGAAGCTTGCATCTCAGCCAGCAGATCTGCTCCCTTCTGGGACCCAGGCTTCAGTGTCACACGGT  ${\tt TCTCAGGCCACCGGCAACTGCATGTACCTCCTCTCTCTCATTTCCAATGATGTACCCTGTACACGTGTTCATG}$  ${\tt TTCCATTTAGATTCAAATGGAGCTAAAATTAAGAGTTTTATGAGCTGTTAAGAATGAGGTAGTTTCTCCTAGGAC}$  $\verb|CCCCCAAAGACAGTGCAAGTAATGACCGTTTGGATCTCATTCGTCGATCTTTGATAGTATGTTCTGGAGTCTACT|\\$ CCCCAGGAGCCAGGCGTGAAGATGGAGTCCTTGTCGCAGCTGGAGCCTTGCCTAGCTGGTGATCACAC  $\verb|CCTGGCCTGTACCTGCACCCCACTGGATGGTGGTACATGGTGGCAGGGACAGGACCACACCCAGTTAAGGCCAGA| \\$ CCAGGCTGAGTGTGACCCCTGAGGTAAACACTCCACTAAGCTGTGTCTTGTTCATGCCCCCTGCTCAGTGAAAGG TGAGTCCCGAGACCAGTTGGGTACCTCTCTATGCGAACCAGAGACATTTCTGGATCCAGGCCAGGTGAAGATTAG GGCCAGGAAGCCTGAGCCCCCGGGGCCTCAAGGTAGGGAGCCGAAGAGGCTGCCAGGACTCTGCTGGGTTGAAAT TTGCCGGGGAGGACTCTTGTCTCCCCCTCAGGAGTATTTTTGTTGAGGCTTTCCTGGAGGTGAAGAAGCAATTCC CATTGCAGCAGGTTAGAGCGAGAATCAGACAGAGGGCAAAAACCAATTCGCTTCTCCCCACGTTCTAAATGCTGG GGCATGGCTGTCAGGAGGGCTTCCTGGGAGGTGTCTCTGGGGGTGAGGTTGGGGCGATGGCCTTTGGAGA TTGCGTGTGGTGTTCAGGACTGTTCCTTGGTGTTTTGAGGGAAACTTTAGTGGGATTGCAGTGGAATGTAAGGTCA

#### 513/2825 FIGURE 461C

## 514/2825 FIGURE 462

MVPPPPGEESQTVILPPGWQSYLSPQGRRYYVNTTTNETTWERPSSSPGIPASPGSHRSSLPPTVNGYHASGTPA HPPETAHMSVRKSTGDSQNLGSSSPSKKQSKENTITINCVTFPHPDTMPEQQLLKPTEWSYCDYFWADKKDPQGN GTVAGFELLLQKQLKGKQMQKEMSEFIRERIKIEEDYAKNLAKLSQNSLASQEEGSLGEAWAQVKKSLADEAEVH LKFSAKLHSEVEKPLMNFRENFKKDMKKCDHHIADLRKQLASRYASVEKARKALTERQRDLEMKTQQLEIKLSNK TEEDIKKARRKSTQAGDDLMRCVDLYNQAQSKWFEEMVTTTLELERLEVERVEMIRQHLCQYTQLRHETDMFNQS TVEPVDQLLRKVDPAKDRELWVREHKTGNIRPVDMEI

#### 515/2825 FIGURE 463

CTGGGCCCCGTGGAACTCCAGCCTGCGCCACCACGTCACGCACACGCTCGGCGCTGCGATCCGCGCATATAACGA  ${\tt TATTTGGATTTGACCTGCATTTTGGAATTTATCTACACTTAAA} \underline{{\tt ATG}} {\tt CCACCAGCAGTTGGAGGTCCAGTTGGATA}$ TCCCAAATCAATTACTGTCTTCTTCAAAGAGATTGAAGGTATATTCCATGCCACCACCAGCGAAGTGTCATGGAT ACTATACGTCTGTATTGGAGTCATTGGAGGTCTTGGGCTTGCCTTCAACTTGAATCCAGCTCTGACCATGATTGG AGCATCCCTTGAGAAAGCTGGAAAATCTGGTGTGAAAAAAGATCTGCATGATGCAAATACAGATCTTATTGGAAG ACACCCTAAACAAGAGAAACGATCAGTCTTCCAAACAATTAATCAGTTCCTGGACTTAACCCTATTCACCCACAG AGCCCGACCATCTATGGGACTTGTAGCCAACACAAAGCCAATAAGACCTCGAATTCAGTATTTCTTTGCGGCTTC CGTTGTTGCAAATGGAGTGTGTCATATGCTAGCACCTTTATCCACTACCTATGTTGGATTCTGTGTCTATGCGGG ATTCTTTGGATTTGCCTTCGGGTGGCTCAGCTCCGTATTGTTTGAAACATTGATGGACCTTGTTGGACCCCAGAG GTTCTCCAGCGCTGTGGGATTGGTGACCATTGTGGAATGCTGTCCTGTCCTCCTGGGGCCCACCACTTTTAGGTCG GCTCAATGACATGTATGGAGACTACAAATACACATACTGGGCATGTGGCGTCGTCCTAATTATTTCAGGTATCTA TCTCTTCATTGGCATGGGCATCAATTATCGACTTTTGGCAAAAGAACAGAAAGCAAACGAGCAGAAAAAGGAAAG TAAAGAGGAAGACCAGTATAGATGTTGCTGGGAAGCCAAATGAAGTTACCAAAGCAGCAGAATCTCCGGACCA GAAAGACACAGAAGGAGGGCCCAAGGAGGAGGAAAGTCCAGTC**TGA**ATCCATGGGGCTGAAGGGTAAATTGAGCA GTTCATGACCCAGGATATCTGAAAATATTCTACTGGCCTGTAATCTACCAGTGGTGCTCAATGCAAATAGTAGAC GAGGGAATGATTGGTAGCAAAGGATGGGGGAAAGAAGTAGGTTCTGTTTTGTTTTTAATCTTAGCTTTTAA TAGTGTCATAAAGATTATAATATGTGCCTTAAGTTTTAGTCTTTAGAACTCTAGAGAGCCTTAACTTCTTAAACC AAAATTTAAAATTTAATCTTGCTTCATTGTTACATGTAATATTTTCAGACATTTTCACTGGAAGATTTATGAACA GAAATATTGGTTGAAAGTTAGAGATTTTACAAAATGCTGACAAAAATATTTTCCTAGCATCAGTAGATTTCTGGC ATATGTTTCTGCTAGCTATATATTTAGGAAATTCAAAGCATAAAACTTTGGCAACATCTTGGCTGTTCTAGACAC AGTGTACTTGTCAACCCCTCTCAGGTACCTTTTCTTGGGATGCTTATTAGAAGCCAAGTAAAGTGCTTAAGGTTT  $\tt GTTTTCATTAAATTAGCTATTTCTGCTCCCCTGTTCAAAGATGCATTTTGAGTGTTTATAGATCACTGCCCTTTT$ TGAAATCACCTGGTATTATTTTTCTTACTGGAAAAGTTAGTATTAAAATCTACAGAACTACATATTTGTGCCTCC TTGGTAAATACAACACTCTAATTAAATGTAGACAGATATTTCAAACATCAGCTGAATTCACTTAAGTTTTTCCA TTTTTTAAGGTTTGCAAATTAAAAAGGGCCAGAAAAATTTGGCACCAGGCAAACGAATAAAGATAGGATTGGGAA AGAAGTTGCTAAGTGTGCTTAGTTTTAATAAGTAATTCCTTCTCTTTTTTCAGAGAAGGCCTTACAGAAAATTGT AAAAGTAGAGAAATGGATTCAGTATAAGGATAAGGAGGGGAAGGTGGACCAGAATGAAAACTGTAAATATTTTTTT AACCTAATATCACTTAAATCGAGGCAGAAAGATATAGACATTCAATGAATTATATTCAATGCATTTAAAATACCA CTGTAATTGACAGAGTAAAAGTATAGATACAAAACCTTGTGTAAGAGGCTGACTTTTCCAAATAAACATTTTTTA AGAAAACATTTCTTCTCCCAAATGTCTATTTTCTTGAGGAAAATATTGCTGTGTCTTCATTTCATTACCAGGTT AAAAAAAAAAAAAAGAAAAAAAAAAAAAAAAG

## 516/2825 FIGURE 464

MPPAVGGPVGYTPPDGGWGWAVVIGAFISIGFSYAFPKSITVFFKEIEGIFHATTSEVSWISSIMLAVMYGGGPI SSILVNKYGSRIVMIVGGCLSGCGLIAASFCNTVQQLYVCIGVIGGLGLAFNLNPALTMIGKYFYKRRPLANGLA MAGSPVFLCTLAPLNQVFFGIFGWRGSFLILGGLLLNCCVAGALMRPĮGPKPTKAGKDKSKASLEKAGKSGVKKD LHDANTDLIGRHPKQEKRSVFQTINQFLDLTLFTHRGFLLYLSGNVIMFFGLFAPLVFLSSYGKSQHYSSEKSAF LLSILAFVDMVARPSMGLVANTKPIRPRIQYFFAASVVANGVCHMLAPLSTTYVGFCVYAGFFGFAFGWLSSVLF ETLMDLVGPQRFSSAVGLVTIVECCPVLLGPPLLGRLNDMYGDYKYTYWACGVVLIISGIYLFIGMGINYRLLAK EQKANEQKKESKEEETSIDVAGKPNEVTKAAESPDQKDTEGGPKEEESPV

#### 517/2825 FIGURE 465

CCCAAATATAAAGGTAGGGAAAGGGACAAGAGGGAACATACCCCTTAGTGTAGAGAAATGGGAAGGAGAAGGAGA AGCCTCAAAAGGAGAGGTGGGAGGGGAATGTCATTAAGGCAGCAAAGTAATCTCTGTAGAAAGATGGAGGAGGAC  $\verb|CCTCCATAGCCTCAGAGATAAAGGCAAAGATTGCCCTCTCAGTGTCCAGAAGGGAAATGGCAGCTTTTCTTCCTT|\\$  ${\tt CCATGGCAGCCACTCCATTGCTCACTCCGGATTACCTTCATCCTTATGTAGATAAGAGTGCTGCAGAGCTCGAAA}$ GGCAGAGATTCGCTTGTGTGGGTTAAAAGTCAGCATTTCCAGCAGCAGCTGTGCTCCCGACTCCTCCATCTCAGG TACCACCGACTGCACTGGGGGGGCCCTCTGGGGGGGAAAGGCTCCACGGGGCAGGGATACATCTCGAGGCCAGTC ATCCTCTGGAGGCAGCCCAATCAGGTCAAAGATTTTGCCCAACTGGTCGGCTTCAGAGTTTCCACAGAAGAGAGG AAGAACTTCGGGAGCTCGGTACCAGAGTGTAACAACCACGGGTGTAAGTGCCATCTGG**TAG**CTGTAGATTCTGGC  ${\tt CAGGCCAAAGTCAGCCAGCTTGACTGTTCCACCACTTGTCACCAGAATGTTCTCTGGCTTCAGATCTCGGTGAACCAGAATGTTCTCTGGCTGAACCAGAATGTTCTCTGGCTTCAGAATCTCTGGGTGAACCAGAATGTTCTCTGGCTTCAGAATCTCTGGTTCAGAATGTTCTCTGGCTTCAGAATGTTCTCTGGCTTCAGAATGTTCTCTGGCTTCAGAATGTTCTAGAATCTCTGGTTCAGAATGTTCTCTGGCTTCAGAATGTTCTCTGGCTTCAGAATGTTCTCTGGCTTCAGAATGTTCTGAGATCTCTGGTTGAACAATGTTCTGAGATCTTCAGAATGTTCTGAGATGTTCTGAAATGTTCTGAGAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCAGAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAAATGTTCTGAATGTTCTGAAATGTTCTGAATGTTCTGAATGTTCTGAATGTTCTGAATGTTAGAATGTTCTGAATGTTAGAATGTTCTGAATGTTAGAATGTTAGAATGTTCTGAATGTAAATGTTCTGAAATGTTAGAATGTAGAATGTTAGAATGTAGAATGTTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGTAGAATGAATGTAGAATGAATGAATGAATGAATGAATGAATGAAATGTAGAATGTAGAATGTAGAATGAAATGAATGAAATGAAATGAATGAATGAATGAAATGAAATGAAATGAAATGAAATGAATGAAATGAA$ GATGCAATTGGCATGAAGGAAATCTAGGCCTCTTAGAAACTGGCGCATCAGATCCTTGATCGTTTCGGCTGGCAA GCCTGGTGGGGGGTGCCTTGTCCAGATATGTCCTTAGGTCCTGGTCTACATGCTCAAACACCAGGGTTACCTTGAT CTCCCGGTCAGTTCGGGATGTGGCACAGACGTCCATCAGCCGGACAACATTGGGATGCTCAAAAGCCTCCAGTCG CCTCAGTAAAGCCACCTCACGAACTGTGCTGATGGGAAGGCCTCCTCCACCTCCTCCATTGGGGACTCTCAC ACTCTTGAGGGCCACAAAGTGGCCACTGTGGGGATCACGGGCCTTGTACACTGTCCCATAGGCACCGACACCAAT TTCAGCCACTGGCTCATATCGAGAGGTAGCCATTCTCAGATCAAGGGAGACCCTCACGCCAGCCCGGGGTGCTGT GGGGGCCCGTTATCGGGGCCCCGGAGCCGGTTCCTACGGCCCCATACACCCGAGCTCGGTCCGGAGCAG

# 518/2825 FIGURE 466

MSTGVAYVDCRRTSGARYQSVTTTGVSAIW

#### 519/2825 FIGURE 467

GAATTCGAGGATCCGGGTACCATGGGAGGAAAACTTCTTCCTGGCCTGGGCTCCGTGCCGCTCTGTTTGCCAACC  $\verb|CTACGACCACCTCTTCAAGTTGCTGATCGGGGGACTCGGGGGGTGGGCAAGACTTGTCTGATCATTCGCTTTGC| \\$ AGAGGACAACTTCAACAACACTTACATCTCCACCATCGGAATTGATTTCAAGATCCGCACTGTGGATATAGAGGG GAAGAAGATCAAACTACAAGTCTGGGACACGGCTGGCCAAGAGCGGTTCAAGACAATAACTACTGCCTACTACCG TGGAGCCATGGGCATTATCCTAGTATACGACATCACGGATGAGAAATCTTTCGAGAATATTCAGAACTGGATGAA AAGCATCAAGGAGAATGCCTCGGCTGGGGTGGAGCCCTCTTGCTGGGGAACAAATGTGACATGGAGGCCAAGAG GAAGGTGCAGAAGGAGCAGGCCGATAAGTTGGCTCGAGAGCATGGAATCCGATTTTTCGAAACTAGTGCTAAATC  $\texttt{AAACGGCAACAAGCCTCCCAGTACTGACCTGAAAACTTGTGACAAGAAGAACACCAACAAGTGCTCCCTGGGC\textbf{\underline{TG}}$ **A**GGACCCTTTCTTGCCTCCCCACCCCGGAAGCTGAACCTGAGGGAGACAACGGCAGAGGGAGTGAGCAGGGGAGA AATAGCAGAGGGCCTTGGAGGGTCACATAGGTAGATGGTAAAGAGAATGAGGAGAAAAAGGAGAAAAGGGAAAAAG GGTAGGAAGAGGGAGGAAAGGAAGGAAGGAGAGACCTTCAGACCTTACCTGGGTTTTCAGGGCA AACATAAATGTAAATACACTGATTTATTCTGTTACTAGATCAGGTTTTAGGGTCCTGCAAAAGGCTAGCTCGGCA AAAAAAAAAAACCATGGTACCCGGATCCTCGAATTC

# 520/2825 FIGURE 468

MAKAYDHLFKLLLIGDSGVGKTCLIIRFAEDNFNNTYISTIGIDFKIRTVDIEGKKIKLQVWDTAGQERFKTITT AYYRGAMGIILVYDITDEKSFENIQNWMKSIKENASAGVERLLLGNKCDMEAKRKVQKEQADKLAREHGIRFFET SAKSSMNVDEAFSSLARDILLKSGGRRSGNGNKPPSTDLKTCDKKNTNKCSLG

### 521/2825 FIGURE 469A

GGTGTGGACGTTGTCTAAATTTCGGTAGCCATGGCACAAGAATATAAGAAAGCATGGGATTATGGCAACCACAGA TGGATTATAACGTTTAGAAGTTCCAATTTTTCAGTGCTTTACAAATAAAGCATCATTTAACCTTTTAAATGAAAA AGATTAAGATCTCATGCAACTGTTGTATTTTCTGGAAGCCATTCTCCAAAAGGGAAGTGCACATTTAAAACACAG ATATGATGGTCCTTGCTGCAGGGATTTAAGTCTACTTGCTTTTACATC**ATG**ACCAGCTTGAAACGGTCACAGACA GAAAGGCCTCTTGCCACTGACAGGGCCTCTGTTGTTGGCACAGACGCCACCCCAAAGTCCACACTGATGATTTC TACATGCGGCGCTTCCGGTCCCAAAATGGCAGCTTAGGATCATCAGTTATGGCTCCTGTAGGACCCCCCGAAGT GAAGGTTCTCACCATATAACCTCAACCCCCGGAGTCCCAAAAATGGGGGTAAGGGCAAGGATTGCAGATTGGCCC CCAAGAAAGGAAAACATAAAAGAATCTAGCCGTTCAAGCCAGGAAATAGAAACCTCAAGTTGCCTTGATAGCCTG TCCTCCAAAAGCAGTCCTGTGAGTCAGGGAAGTTCTGTTAGCCTCAATTCCAATGACTCAGCCATGCTGAAAAGC ATACAGAACACGCTGAAAAACAAGACAAGACCGTCGGAGAACATGGACTCCAGATTTCTCATGCCTGAAGCCTAC CCCAGCTCCCCCAGAAAAGCTCTTCGCAGAATACGCCAGCGAAGCAACAGTGATATCACCATAAGTGAACTTGAT GTGGATAGCTTTGATGAATGTATCTCACCTACATACAAGACTGGACCATCACTGCACAGGGAATATGGTAGCACA TCTTCAATTGATAAACAGGGAACATCTGGAGAAAGCTTTTTTGATTTGATAAGGGCTACAAAGATGACAAATCT GATCGAGGTCCAACTCCAACCAAGCTCAGTGACTTTCTCATTACTGGTGGTGGCAAGGGTTCTGGTTTCTCTTTG GATGTAATAGACGGGCCTATCTCACAGAGAGAGACCTCAGGCTTTTTAAGGAAAAGGGAAAAACCACTCAAGCGA CGTTCAAAATCTGAAACTGGAGACTCCTCTATTTTTCGTAAATTGCGCAATGCCAAAGGTGAAGAACTTGGGAAG TCATCAGATCTTGAAGATAACCGATCAGAAGACTCTGTCAGGCCCTGGACATGTCCAAAGTGCTTTGCCCACTAT GATGTCCAGAGTATATTATTTGATTGAATGAGGCAATTATGAACAGGCACAATGTTATTAAGAGGAGAAACACC ACCACTGGAGCTTCCGCAGCTGCCGTGGCATCCTTGGTCTCTGGACCTCTGTCTCATTCAGCCAGTTTTAGCTCC CCAATGGGCAGCACAGAGGACCTGAATTCCAAAGGAAGCCTCAGCATGGACCAGGGAGATGATAAAAGCAATGAG AATTCTGGCTCCTTTAGTGGATGTGAAAGTGCCTCCTTTGAGTCTACCCTTAGTTCCCATTGCACAAATGCAGGA GTAGATCTGGGTGCATACTATTATAGAAAATTTTTCTACCAGAAGGAACACTGGAACTATTTTGGGGCTGATGAG AATCTTGGTCCAGTGGCTGTGAGCATTCGAAGGGAAAAACCAGATGAAATGAAAAAATGGATCTCCGTACAAC TACCGAATAATTTTTAGAACTAGTGAGCTCATGACACTGAGAGGTTCGGTCCTGGAGGACGCCATTCCGTCGACA GCCAAGCACTCGACAGCCAGAGGCCTGCCTCTCAAAGAAGTGCTGGAGCACGTGGTTCCTGAGCTCAATGTCCAG TGCCTGCGGTTGGCCTTCAACACACCCAAGGTCACAGAGCAGCTCATGAAACTGGATGAACAAGGGCTGAACTAC CAGCAGAAAGTAGGCATCATGTACTGCAAAGCTGGACAGAGCACTGAAGAAGAGATGTACAACAATGAGTCAGCT GGCCCAGCCTTTGAAGAATTCCTTCAACTATTGGGAGAGCGAGTTCGGCTCAAAGGATTTGAGAAGTATCGAGCA CAGCTTGATACCAAAACTGACTCCACTGGAACCCATTCTCTGTACACAACATACAAAGATTATGAAATTATGTTC CATGTTTCTACCATGCTGCCATACACCCCAACAACAACAACAGCTCCTGAGGAAGCGGCACATTGGAAATGAT ATCGTAACAATTGTTTTCCAAGAGCCTGGAGCACAGCCATTCAGCCCAAAAAACATCCGATCCCACTTCCAGCAC GTTTTCGTCATCGTCAGGGTGCACAATCCGTGCTCTGACAGTGTCTGTTATAGTGTGGCTGTTACCAGGTCCAGA GATGTGCCTTCCTTTGGGCCTCCCATTCCTAAAGGGGTCACTTTCCCTAAGTCAAATGTGTTCAGGGACTTCCTT  $\tt TTGGCGAAAGTGATTAATGCAGAAAATGCTGCTCATAAATCGGAGAAGTTTCGGGCCATGGCAACTCGGACCCGC$ CAGGAATACCTGAAAGATCTGGCAGAAAAGAATGTCACCAACACCCCTATCGACCCTTCTGGCAAGTTTCCGTTC ATCTCTCTGGCTTCCAAGAAGAAGGAAAAGTCTAAGCCATATCCAGGAGCCGAGCTCAGCAGCATGGGGGCCATT GTATGGGCAGTCCGGGCTGAAGACTACAACAAGGCCATGGAACTAGACTGCCTTTTAGGGATCTCCAATGAGTTC ATTGTGCTCATTGAACAGGAAACAAGAGCGTGGTCTTCAATTGTTCCTGTAGAGATGTGATAGGGTGGACTTCA ACTGACACCAGCCTCAAAATCTTCTATGAACGAGGAGAATGTTTTCAGTGGGTAGTTTTATTAACATTGAGGAG ATCAAAGAGATTGTCAAAAGGTTGCAGTTTGTTTCAAAAGGCTGTGAATCGGTGGAGATGACTCTGCGAAGAAAT GGGCTAGGACAGCTTGGCTTCCATGTCAACTATGAGGGCATTGTGGCGGATGTGGAGCCCTACGGTTATGCCTGG CAGGCAGGGCTGAGGCAGTCGCCTGGTGGAGATCTGCAAGGTGGCGGTAGCCACTCTGAGCCATGAGCAG ATGATCGACCTCCTGAGAACATCTGTCACGGTGAAGGTTGTCATCATTCCCCCGCATGATGACTGCACCCCGCGG TTTCCCTTCCGAAATAATAACAAGTGGCAGAGGAACGCCAGCAAGGGGCCTCATTCACCTCAAGTCCCGTCCCAG

#### 522/2825 FIGURE 469B

GTGCAGAGTCCCATGACCTCGCGGCTGAATGCTGGAAAAGGAGATGGGAAGATGCCTCCTCCAGAAAGAGCCGCC AACATCCCTCGAAGCATCTCCAGTGACGGCCCCACTAGAGAGGCGGCTGTCTCCTGGTTCGGACATCTATGTG ACGGTCTCATCCATGGCTTTAGCAAGATCCCAGTGTCGGAACTCTCCTAGCAACTTGTCTTCATCCAGTGATACT GGTTCTGTGGGGGGCACTTACAGGCAGAAGTCCATGCCCGAAGGGTTTGGAGTGAGCCGTAGATCCCCAGCCTCC ATTGACAGGCAGAACACCCAGTCAGATATTGGTGGCAGCGGAAAATCCACGCCTAGCTGGCAAAGAAGTGAGGAT AGCATTGCTGACCAGATGGCTTACAGTTATAGAGGACCTCAGGATTTCAATTCTTTTGTCCTCGAGCAGCATGAA TTAATGCGGCAGGATCCAGTGGTTCATTTGTCTCCAAACAACAAGGGCATTCTGATAGCCACTACTCGAGCCAC TCCAGTAGCAATACTCTCCCAGCAATGCGTCAAGTGCCCATAGTGATGAGAAGTGGTACGATGGGGACCGCACA GAATCCGAACTCAACAGCTATAACTATCTGCAAGGCACCTCTGCTGACAGTGGCATTGACACCACCTCTTATGGC CCCAGCCACGCAGCACACCTCGCTGGGGGCTGCCACATCGTCACCTCGCTCAGGGCCAGGCAAGGAGAAAGTG GCACCCCTATGGCACAGCTCCAGTGAAGTAATCTCCATGGCAGATCGGACTTTGGAGACAGAGAGCCACGGCCTG GACCGGAAAACAGAGTCTTCCCTGAGCTTAGACATACACAGCAAGAGCCAAGCCGGCTCGACCCCTCTGACAAGG GAGAACAGCACCTTCAGTATAAACGATGCTGCTTCCCACACAAGTACCATGAGCTCCCGACACTCTGCCAGCCCA GTGGTTTTCACCAGTGCCCGGAGTTCACCTAAAGAAGAGCTTCATCCAGCTGCCCCCTCACAGCTCGCACCATCC TTCTCCTCCTCCTCCTCCTCTGGTCCTAGGAGTTTTTACCCTCGCCAGGGCGCTACTAGCAAGTACCTG ATTGGATGGAAAAACCCGAAGGAACCATAAACTCCGTGGGATTTATGGACACGAGAAAGCGTCATCAGAGCGAT GGCAATGAAATAGCCCACACCAGGCTGCGTGCCTCAACCAGAGACCTCCGGGCATCTCCTAAGCCAACCTCCAAG TCCACCATTGAAGAAGATCTAAAGAAACTAATTGATCTTGAAAGCCCAACTCCTGAATCACAGAAGAGTTTTAAG TTCCACGCACTCTCCTCAGTCTCCTTTCCCCAGCACCCCCACCTCACGGCGGGCCTTGCACAGAACACTG TCGGACGAGAGCATTTACAATAGCCAGAGGGAGCACTTTTTCACCTCCAGGGCGTCACTTCTGGACCAAGCCCTG CCCAACGACGTCCTCTTCAGTAGCACGTACCCTTCTCTCCCCAAGTCGCTCCCGTTGAGGAGGCCTTCTTACACC TTAGGAATGAAATCGCTGCATGGAGAGTTCTCAGCCTCGGACAGCTCCCTCACTGACATCCAGGAGACCCGCAGG CAGCCTATGCCCGACCCTGGCCTGATGCCCTGCCTGACACTGCTGCAGACTTGGATTGGTCCAACCTGGTAGAT GCTGCCAAAGCCTATGAGGTCCAGAGAGCCTCATTTTTTGCTGCTAGTGATGAAAACCATCGCCCCTTGAGTGCT GCATCCAACAGTGATCAGCTGGAGGACCAGGCTCTGGCCCAGATGAAGCCTTACAGCAGCAGTAAAGACTCCTCT CCCACTCTGGCTTCTAAAGTGGACCAGCTGGAAGGTATGCTGAAGATGCTTCGGGAAGATTTTGAAGAAGGAAAAA GAAGACAAAGCTCACCTTCAGGCGGAGGTGCAGCACCTGCGAGAGGACAACCTGAGGCTACAGGAGGAGTCCCAG AACGCCTCGGACAAGCTGAAGAAGTTCACAGAATGGGTCTTCAACACCATAGACATGAGCTAGGGAAGGCTGAGG AGGACAGGAGAAGGGCCCAGACACTCCCTCCAGTGAGTGTCCTGCAGCCCTTATTCCCTCCATAGAAAGCATCCT CAGAGCACCTTCCCTGGCTTCCTACTCTGCCCCCTTTCGGGGAGTGCACAACACAATAGTTGCAGATCAACAATC ATCACCTGCCTTTTGTAGAAAAGAAAAACAAAAAAGTAAAAATTTTTAAACAGTAAAATAAAAGTTTAACT GCTAAAAAAAAAAAAAAAAAAAAAA

#### 523/2825 FIGURE 470

MTSLKRSQTERPLATDRASVVGTDGTPKVHTDDFYMRRFRSQNGSLGSSVMAPVGPPRSEGSHHITSTPGVPKMG VRARIADWPPRKENIKESSRSSQEIETSSCLDSLSSKSSPVSQGSSVSLNSNDSAMLKSIQNTLKNKTRPSENMD  ${\tt SRFLMPEAYPSSPRKALRRIRQRSNSDITISELDVDSFDECISPTYKTGPSLHREYGSTSSIDKQGTSGESFFDL}$ LKGYKDDKSDRGPTPTKLSDFLITGGGKGSGFSLDVIDGPISQRENLRLFKEREKPLKRRSKSETGDSSIFRKLR NAKGEELGKSSDLEDNRSEDSVRPWTCPKCFAHYDVQSILFDLNEAIMNRHNVIKRRNTTTGASAAAVASLVSGP LSHSASFSSPMGSTEDLNSKGSLSMDQGDDKSNELVMSCPYFRNEIGGEGERKISLSKSNSGSFSGCESASFEST LSSHCTNAGVAVLEVPKENLVLHLDRVKRYIVEHVDLGAYYYRKFFYQKEHWNYFGADENLGPVAVSIRREKPDE MKENGSPYNYRIIFRTSELMTLRGSVLEDAIPSTAKHSTARGLPLKEVLEHVVPELNVQCLRLAFNTPKVTEQLM KLDEQGLNYQQKVGIMYCKAGQSTEEEMYNNESAGPAFEEFLQLLGERVRLKGFEKYRAQLDTKTDSTGTHSLYT TYKDYEIMFHVSTMLPYTPNNKQQLLRKRHIGNDIVTIVFQEPGAQPFSPKNIRSHFQHVFVIVRVHNPCSDSVC YSVAVTRSRDVPSFGPPIPKGVTFPKSNVFRDFLLAKVINAENAAHKSEKFRAMATRTRQEYLKDLAEKNVTNTP IDPSGKFPFISLASKKKEKSKPYPGAELSSMGAIVWAVRAEDYNKAMELDCLLGISNEFIVLIEQETKSVVFNCS CRDVIGWTSTDTSLKIFYERGECVSVGSFINIEEIKEIVKRLQFVSKGCESVEMTLRRNGLGQLGFHVNYEGIVA DVEPYGYAWQAGLRQGSRLVEICKVAVATLSHEQMIDLLRTSVTVKVVIIPPHDDCTPRRSCSETYRMPVMEYKM NEGVSYEFKFPFRNNNKWQRNASKGPHSPQVPSQVQSPMTSRLNAGKGDGKMPPPERAANIPRSISSDGRPLERR  ${\tt TPSWQRSEDSIADQMAYSYRGPQDFNSFVLEQHEYTEPTCHLPAVSKVLPAFRESPSGRLMRQDPVVHLSPNKQG}\\$ HSDSHYSSHSSSNTLSSNASSAHSDEKWYDGDRTESELNSYNYLQGTSADSGIDTTSYGPSHGSTASLGAATSSP RSGPGKEKVAPLWHSSSEVISMADRTLETESHGLDRKTESSLSLDIHSKSQAGSTPLTRENSTFSINDAASHTST  $ext{MSSRHSASPVVFTSARSSPKEELHPAAPSQLAPSFSSSSSSSSGPRSFYPRQGATSKYLIGWKKPEGTINSVGFM}$ DTRKRHQSDGNEIAHTRLRASTRDLRASPKPTSKSTIEEDLKKLIDLESPTPESQKSFKFHALSSPQSPFPSTPT SRRALHRTLSDESIYNSQREHFFTSRASLLDQALPNDVLFSSTYPSLPKSLPLRRPSYTLGMKSLHGEFSASDSS  ${ t LTDIQETRRQPMPDPGLMPLPDTAADLDWSNLVDAAKAYEVQRASFFAASDENHRPLSAASNSDQLEDQALAQMK}$ PYSSSKDSSPTLASKVDQLEGMLKMLREDLKKEKEDKAHLQAEVQHLREDNLRLQEESQNASDKLKKFTEWVFNT TDMS

## 524/2825 FIGURE 471

 $\tt CGGAGGTGAGGTTTGTTACCGCGATTCTGAGAGGTGGGCTTTTAGTCCCTCCAGACCTCGGCTTTAGTGCTGTCT$ CCGCTTTTCTTTCACCTTCACAGAGGTTCGTGTCTTCCTAAAAGAAGGTTTTATTGGGAGGTAAAGGTCAATGCG TAGGGGTAGAGTAAG**ATG**TCTTATGGTGAAATTGAAGGTAAATTCTTGGGACCTAGAGAAGAAGTAACGAGTGAG CCACGCTGTAAAAAATTGAAGTCAACCACAGAGTCGTATGTTTTTCACAATCATAGTAATGCTGATTTTCACAGA ATCCAAGAGAAAACTGGAAATGATTGGGTCCCTGTGACCATCATTGATGTCAGAGGACATAGTTATTTGCAGGAG AACAAAATCAAAACTACAGATTTGCATAGACCTTTGCATGATGAGACCTGGTAATAGACCAGATGTTATTGAA TCCATTGATTCACAGGTTTTACAGGAAGCACGTCCTCCATTAGTATCCGCAGACGATGAGATATATAGCACAAGT AAAGCATTTATAGGACCCATTTACAAACCCCCTGAGAAAAAGAAACGTAATGAAGGGAGGAATGAGGCACATGTT CTAAATGGTATAAATGACAGAGGAGGACAAAAAGAGAAACAGAAATTTAACTCTGAAAAATCAGAGATTGACAAT GAATTATTCCAGTTTTACAAAGAAATTGAAGAGCTTGAAAAGGAAAAAGATGGTTTTGAGAACAGTTGTAAAGAA TCTGAACCTTCTCAGGAACAATTTGTTCCATTTTATGAGGGTCATAATAATGGTCTCTTAAAACCTGATGAAGAA AAGAAAGATCTTAGTAATAAAGCTATGCCATCACATTGTGATTATCAGCAGAACTTGGGGAATGAGCCAGACAAA TATCCCTGTAATGGACAAGTAATACCTACATTTTGTGACACTTCATTTACTTCTTTCAGGCCTGAATGGCAGTCA GTATATCCTTTTATAGTGCCCTATGGTCCCCCTCTTCCCAGTTTGAACTATCATTTAAACATTCAGAGATTCAGT GGTCCACCAAATCCACCATCAAATATTTTCCAAGCCCAAGATGACTCTCAGATACAAAATGGATATTATGTAAAT AGTGTTCATCCCTCTGGAAATGGCTGCAGTATGCAAGATCGATATGTGAGTAATGGTTTCTGTGAAGTCAGAGAA AAGTTAAATAAATTGCAGAAGTTACTTATTCTTTTAAGAGGTCTGCCTGGTTCTGGGAAAACAACATTGTCTCGA ATTCTGCTTGGTCAGAATCGTGATGGCATTGTGTTCAGCACTGATGACTATTTTCACCATCAAGATGGGTACAGG TATAATGTTAATCAACTTGGTGATGCCCATGACTGGAACCAGAACAGAGCAAAACAAGCTATCGATCAGGGAAGA TCTCCAGTTATAATAGATAACACTAATATACAAGCTTGGGAAATGAAGCCATATGTGGAAGTGGCCATAGGAAAA GGATACAGAGTAGAGTTTCATGAACCTGAAACTTGGTGGAAATTTGATCCTGAAGAATTAGAAAAGAGGAATAAA CATGGTGTGTCTCGAAAGAAGATTGCTCAGATGTTGGATCGTTATGAATATCAAATGTCCATTTCTATTGTAATG AATTCAGTGGAACCATCACACAAAAGCACACAAAGACCTCCTCCTCCACAGGGGAACAGAGGTGGGGAGGCTCT  $\verb|CTTGGCTCACATAATCGTGTCTGTGTCACAAATAATCATTAAATTAGCTATTTTCAGCTAACACATTTGTTGTTG| \\$ CACTTGAAAAAGAGTTAGTGAGCCTGTCTTGGAGTTTAAGTAGTTTCAAATAAAAAAAGGCTACAGTGCCTCACA TTAAATGTTTTTATATTCTCTTGTTGTAATACTCTTGGCTGTTATGGAAGCACCTGAGTAATAGAGTGGTGGGTA GGAGCTAGGATGTTTTTCTACAATCGAATTTTAAACTAATTTATCTATTTTATAGACACTATTGAACAGTTTTTT AATAGTTCATATCTAAATCTAACTTTTCATAAAACTTTACGGTTTTTCCTTCACTACCTTAAATATGCAAGAAAT CCTCCTTCTCAACCCATCTCCCTAACCCCACATGCTTGCCAGTTCTTGAAGGGATAAAGTGATTCTAATAATGTT .TTACTTCTCTGTTCAATTTAATGTGATATAATTCTAGTATAAAAATATTTTGGACAGTTGCTTAACATGGTCA TAAGAGGATTTGTACTATAGAATATCTTCTAGTACTAATTTTTCTGTAGAGCAAATTATATTTCTCTCACTGGAT AGTTTTTAGATGTGTTTCTTCATATAAAATTAAAAACTGAGATGGAATTC

## 525/2825 FIGURE 472

MSYGEIEGKFLGPREEVTSEPRCKKLKSTTESYVFHNHSNADFHRIQEKTGNDWVPVTIIDVRGHSYLQENKIKT TDLHRPLHDEMPGNRPDVIESIDSQVLQEARPPLVSADDEIYSTSKAFIGPIYKPPEKKKRNEGRNEAHVLNGIN DRGGQKEKQKFNSEKSEIDNELFQFYKEIEELEKEKDGFENSCKESEPSQEQFVPFYEGHNNGLLKPDEEKKDLS NKAMPSHCDYQQNLGNEPDKYPCNGQVIPTFCDTSFTSFRPEWQSVYPFIVPYGPPLPSLNYHLNIQRFSGPPNP PSNIFQAQDDSQIQNGYYVNNCHVNWNCMTFDQNNEYTDCSENRSSVHPSGNGCSMQDRYVSNGFCEVRERCWKD HCMDKHNGTDRFVNQQFQEEKLNKLQKLLILLRGLPGSGKTTLSRILLGQNRDGIVFSTDDYFHHQDGYRYNVNQ LGDAHDWNQNRAKQAIDQGRSPVIIDNTNIQAWEMKPYVEVAIGKGYRVEFHEPETWWKFDPEELEKRNKHGVSR KKIAQMLDRYEYQMSISIVMNSVEPSHKSTQRPPPPQGRQRWGGSLGSHNRVCVTNNH

#### 526/2825 FIGURE 473

GGCGGCGGCAGCGGTTCGGTTGCGCGTGGCGCACGGGTGGGAGCCGAGCCCAGGCCGGAGCAGGCGCCCCC GTGGGGCTGCCGCCCTCGCCCCCCGGGCTTCACGCTGCCGCCGCTGGGAGGCAGCCTGGGCCCGGC ACCAGTACGAGTCGAAGTTCGGAACGGACCCCCGGGGCTGCAACCGCCAGCGCCTCAGGGGCCGCCGAGGATGGG GCCTGCGGCTGCCCAACCCGGGCACATTCGAGGAGTGCCACCGGAAGTGCAAGGAGCTGTTTCCCATTCAG ATGGAGGGTGTCAAGCTCAACAAAGGGTTGAGTAACCATTTTCAGGTCAACCACACAGTAGCCCTCAGC ACAATCGGGGAGTCCAACTACCACTTCGGGGTCACATATGTGGGGACAAAGCAGCTGAGTCCCACAGAGGCGTTC CCTGTACTGGTGGGTGACATGGACAACAGTGGCAGTCTCAACGCTCAGGTCATTCACCAGCTGGGCCCCGGTCTC AGGTCCAAGATGGCCATCCAGACCCAGCAGTCGAAGTTTGTGAACTGGCAGGTGGACGGGGAGTATCGGGGCTCT GACTTCACAGCAGCCGTCACCCTGGGGAACCCAGACGTCCTCGTGGGTTCAGGAATCCTCGTAGCCCACTACCTC CAGAGCATCACGCCTTGCCTGGCCCTGGGTGGAGAGCTGGTCTACCACCGGCGGCCTGGAGAGGGGGCACTGTC ATGTCTCTAGCTGGGAAATACACATTGAACAACTGGTTGGCAACGGTAACGTTGGGCCAGGCGGCATGCACGCA AGCGTCTCCTTCGGGTACCAGCTGGACCTGCCCAAGGCCAACCTCCTCTTCAAAGGCTCTGTGGATAGCAACTGG ATCGTGGGTGCCACGCTGGAGAAGAAGCTCCCACCCCTGCCCCTGACACTGGCCCTTGGGGCCTTCCTGAATCAC  $\tt CGCAAGAACAAGTTTCAGTGTGGCCTTTGGCCTCACCATCGGCT{\tt CACCCCTT}{\tt CACCCCCTT}{\tt CACCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCCTT}{\tt CACCCCTT}{\tt CACCCTT}{\tt CACCCCTT}{\tt CACCCTT}{\tt CACCTT}{\tt C$ GGGGGTCGGGGGGGACATTGGAAAGGAGGGACCCCGCCACCCCAGCAGCTGAGGAGGGGATTCTGGAACTGAAT GGGGCACGCAGGATTCTGAGCACCAGGGGCAGAGGCGCCAGACAACCTCAGGGAGGAGTGTCCTGGCGTCCCCA TCCTCCAAAGGGCCTGGGCCCCGAGGGGGGCAGCGAGAGGAGCTTCCCCATCCCCGGTCAGTCCACCCTGCC  ${\tt CCGTCCACTTTCCCATCTCCTCGGTATAAATCATGTTTATAAGTTATGGAAGAACCGGGACATTTTACAGAAAAA}$ 

## 527/2825 **FIGURE 474**

MGNVLAASSPPAGPPPPPAPALVGLPPPPPPSPPGFTLPPLGGSLGAGTSTSRSSERTPGAATASASGAAEDGACG CLPNPGTFEECHRKCKELFPIQMEGVKLTVNKGLSNHFQVNHTVALSTIGESNYHFGVTYVGTKQLSPTEAFPVL VGDMDNSGSLNAQVIHQLGPGLRSKMAIQTQQSKFVNWQVDGEYRGSDFTAAVTLGNPDVLVGSGILVAHYLQSI TPCLALGGELVYHRRPGEEGTVMSLAGKYTLNNWLATVTLGQAGMHATYYHKASDQLQVGVEFEASTRMQDTSVS FGYQLDLPKANLLFKGSVDSNWIVGATLEKKLPPLPLTLALGAFLNHRKNKFQCGFGLTIG

#### 528/2825 FIGURE 475

GCAGAGGCGCAGGTAGATGGAGTTGGGGAGTTGCCTGGAGGGCGGAGGAGGAGGCGGAGGAGGAGAGAGGGCGAGCC TGAGGTGAAAAAGCGGCGACTTCTGTGTGTGAGGTTTGCCTCGGTCGCAAGCTGCGATGCCGCAGTGGCTCAGTG CTTGGAACGCCGACCTGAAACCATCTCTGAGCCCAAGACCTATGTTGACCTAACCAATGAAGAAACAACTGATTC TATTGATGGATTAGATCTAAACAATCTGTCAGAGGGGCTCGAGGGGTGTGTTCCTACTTAGCTTTGTACAGCCC AGATGTGATATTTCTACAGGAAGTTATTCCCCCCATATTATAGCTACCTAAAGAAGAGATCAAGTAATTATGAGAT TATTACAGGTCATGAAGAAGGATATTTCACAGCTATAATGTTGAAGAAATCAAGAGTGAAATTAAAAAGCCAAGA GATTATTCCTTTTCCAAGTACCAAAATGATGAGAAACCTTTTATGTGTGCATGTGAACGTGTCAGGAAATGAGCT AAAGAAAATGCAAGAGGCTCCAGAGTCAGCTACAGTTATATTTGCAGGAGATACAAATCTAAGGGATCGAGAGGT TACCAGATGTGGTGGTTTACCCAACAACATTGTGGATGTCTGGGAGTTTTTGGGCAAACCTAAACATTGCCAGTA TACATGGGATACACAAATGAACTCTAATCTTGGAATAACTGCTGCTTGTAAACTTCGTTTTGATCGAATATTTTT CAGAGCAGCAGCAGAAGAGGGACACATTATTCCCCGAAGTTTGGACCTTCTTGGATTAGAAAAACTGGACTGTGG  $\texttt{TAGATITCCTAGTGATCACTGGGGTCTTCTGTGCAACTTAGATATAATATTG} \underline{\textbf{TAA}} \texttt{AATGCTTTTCAAGTGTGGGT}$ ATCATTGTCACAGAAAAACCAACTATGATTTATGGTTGTGTTTTCAGAATTCAACATTAAAGATTAATGTTTATT TAAACGAACACATTCCTGCATTCAGGATGTGAGGCCATTTAATAAAAAGGGCACAAAGCCTGTCAGAGTTTTCAA CGGTGCTTATAGCTGCCAGCTGGATTCCAAACAGGTACCCCATTGTCTCTGAGCTAATGTTTATATTTTTCCATT CAGGCACCGAAATAGTTAATATTTAAAATAAGTCTTCAAAAGAAAACATAAGAGATTATTGAGTTCTTGGGACTG GATCCTTTATTTCATAAGTTCAGATCATCTTAAATGAAAATGCCATGATTATCTGCAGTTAAGTAGATGACAGCT ATTCTACATCAGACTTGATTTTTGTCAGCTAATTACATAATTGGTAAGCTATAATTGAAACCTTATGGCTTAAAA TTCCTTAACTCCTTTTTGATTCATGTTTGTAGTCATGTTGTCAACAGAGGCAAAGTTAAGCTTGATGATGGTTAA CTAATTGTGAAATGGAAATGCTTTACAGGAAGTAAATGCAAATTACTTTTAAGTGTGCTTTAAAGAAAAATATTT 

## 529/2825 FIGURE 476

MELGSCLEGGREAAEEEGEPEVKKRRLLCVRFASVASCDAAVAQCFLAENDWEMERALNSYFEPPVEESALERRP ETISEPKTYVDLTNEETTDSTTSKISPSEDTQQENGSMFSLITWNIDGLDLNNLSERARGVCSYLALYSPDVIFL QEVIPPYYSYLKKRSSNYEIITGHEEGYFTAIMLKKSRVKLKSQEIIPFPSTKMMRNLLCVHVNVSGNELCLMTS HLESTRGHAAERMNQLKMVLKKMQEAPESATVIFAGDTNLRDREVTRCGGLPNNIVDVWEFLGKPKHCQYTWDTQ MNSNLGITAACKLRFDRIFFRAAAEEGHIIPRSLDLLGLEKLDCGRFPSDHWGLLCNLDIIL

#### 530/2825 FIGURE 477

GCTGGGAAAGCTGCTCAAGGAGCAGAAGTACGACCGGCAGCTGAGGTTGTGGGGTGATCATGGGCAAGAGGCTTT AGAATCTGCTCATGTTTGCCTAATAAATGCAACAGCCACAGGAACTGAAATTCTTAAAAACTTGGTACTACCAGG AAGCAGTATCGGCAAGAACCGAGCTGAAGCTGCCATGGAATTCTTACAAGAATTAAATAGCGATGTCTCTGGAAG  $\verb|TTTTGTGGAAGAGTCCAGAAAACCTTCTAGACAATGATCCCTCATTTTTCTGTAGGTTTACTGTTGTAGTTGC|\\$ AACTCAGCTTCCTGAAAGCACTTCACTACGCTTAGCAGATGTCCTCTGGAATTCCCAGATTCCTCTTTTGATCTG TAGGACATATGGACTAGTTGGTTATATGAGGATCATTATAAAAGAACATCCAGTAATAGAATCTCATCCAGATAA TGCATTAGAGGATCTACGACTAGATAAGCCATTTCCTGAACTGAGAGAACATTTTCAGTCCTATGATTTGGATCA TATGGAAAAAAGGACCACAGTCATACTCCATGGATTGTGATCATAGCTAAATATTTAGCACAGTGGTATAGTGA AACAAATGGACGAATACCTAAAACGTATAAAGAAAAAGGGACTTCAGAGATTTGATTAGACAAGGAATTCTAAA AAATGAAAATGGGGCTCCAGAAGATGAAGAGAATTTTGAAGAAGCTATTAAAAATGTGAACACAGCACTAAATAC AACTCAGATCCCAAGCAGTATTGAAGATATATTTAATGATGATCGCTGCATAAATATCACCAAACAGACTCCATC ATTTTGGATTTTAGCTCGTGCCTTAAAGGAATTTGTGGCCAAAGAGGGTCAAGGAAATTTACCTGTTCGAGGCAC TGCTGCCGCTGTGGGTAATCATGTTGCCAAATTGCTGCAGTCCATTGGCCAGGCACCAGAGTCCATTTCAGAGAA AGAATTAAAATTACTCTGCAGCAATTCTGCATTTCTTCGAGTGGTAAGATGTCGATCCTTAGCTGAAGAATATGG TTTGGATACAATTAACAAGGATGAAATTATTTCTAGCATGGACAATCCAGATAATGAAATAGTGTTGTACTTAAT GTTACGGGCTGTTGATAGATTTCATAAACAACAGGGTAGATATCCAGGAGTATCTAACTATCAAGTTGAAGAAGA TATAGGAAAGTTGAAGTCTTGTCTCACTGGCTTCCTTCAGGAATATGGTTTATCTGTAATGGTGAAAGATGATTA AGAGGTCATCAAAATAATCACCAAACAATTTGTAATTTTTAATAATACTTACATTTACAGTGGCATGTCACAAAC TTCAGCAACTTTCCAGTTG**TAG**AGTAAGCAAGCACCTTAAGTAGTGTGTTAATGATTGAAACTGTAATTGCCTTC GGGTTGTGCTTTAGTCTGTAAAATTCTAAAGGAGAGCTGCTAAATTGTTTTCTTAATAAACATTTTTCTCATTTG TAAAAAAAA

# 531/2825 FIGURE 478

MAQLGKLLKEQKYDRQLRLWGDHGQEALESAHVCLINATATGTEILKNLVLPGIGSFTIIDGNQVSGEDAGNNFF LQRSSIGKNRAEAAMEFLQELNSDVSGSFVEESPENLLDNDPSFFCRFTVVVATQLPESTSLRLADVLWNSQIPL LICRTYGLVGYMRIIIKEHPVIESHPDNALEDLRLDKPFPELREHFQSYDLDHMEKKDHSHTPWIVIIAKYLAQW YSETNGRIPKTYKEKEDFRDLIRQGILKNENGAPEDEENFEEAIKNVNTALNTTQIPSSIEDIFNDDRCINITKQ TPSFWILARALKEFVAKEGQGNLPVRGTIPDMIADSGKYIKLQNVYREKAKKDAAAVGNHVAKLLQSIGQAPESI SEKELKLLCSNSAFLRVVRCRSLAEEYGLDTINKDEIISSMDNPDNEIVLYLMLRAVDRFHKQQGRYPGVSNYQV EEDIGKLKSCLTGFLQEYGLSVMVKDDYVHEFCRYGAAEPHTIAAFLGGAAAQEVIKIITKQFVIFNNTYIYSGM SQTSATFQL

### 532/2825 FIGURE 479

AGGGACAGCCCAGAGGAGGCGTGGCCACGCTGCCGGCGGAAGTGGAGCCCTCCGCGAGCGCGCGAGGCCGCGGG GCAGGTGCCCGAGAGGCAGGGGCTGGCCTGGGATGCGCGCACCTGCCCTCGCCCCGCCCCGCACGAGGG GCCGATTAAATGGGCCGGCGGGCTCAGCCCCCGGAAACGGTCGTAACTTCGGGGCTGCGAGCGCGGAGGGCGAC GACGACGAAGCGCAGACAGCGTC<u>ATG</u>GCAGAGCAGGTGGCCCTGAGCCCGGACCCAGGTGTGCGGGATCCTGCGGG AAGAGCTTTTCCAGGGCGATGCCTTCCATCAGTCGGATACACACATATTCATCATCATGGGTGCATCGGGTGACC TGGCCAAGAAGAAGATCTACCCCACCATCTGGTGGCTGTTCCGGGATGGCCTTCTGCCCGAAAACACCTTCATCG  ${\tt TGGGCTATGCCCGTTCCCGCCTCACAGTGGCTGACATCCGCAAACAGAGTGAGCCCTTCTTCAAGGCCACCCCAG}$ ACCAGCGCCTCAACAGCCACATGGATGCCCTCCACCTGGGGTCACAGGCCAACCGCCTCTTCTACCTGGCCTTGC CCCCGACCGTCTACGAGGCCGTCACCAAGAACATTCACGAGTCCTGCATGAGCCAGATAGGCTGGAACCGCATCA TCGTGGAGAGCCCTTCGGGAGGGACCTGCAGAGCTCTGACCGGCTGTCCAACCACATCTCCCTGTTCCGTG AGGACCAGATCTACCGCATCGACCACTACCTGGGCAAGGAGATGGTGCAGAACCTCATGGTGCTGAGATTTGCCA ACAGGATCTTCGGCCCCATCTGGAACCGGGACAACATCGCCTGCGTTATCCTCACCTTCAAGGAGCCCTTTGGCA CTGAGGGTCGCGGGGCTATTTCGATGAATTTGGGATCATCCGGGACGTGATGCAGAACCACCTACTGCAGATGC TGTGTCTGGTGGCCATGGAGAAGCCCGCCTCCACCAACTCAGATGACGTCCGTGATGAGAAGGTCAAGGTGTTGA AATGCATCTCAGAGGTGCAGGCCAACAATGTGGTCCTGGGCCAGTACGTGGGGAACCCCGATGGAGAGGGCGAGG CCACCAAAGGGTACCTGGACGACCCCACGGTGCCCCGCGGGTCCACCACCGCCACTTTTGCAGCCGTCGTCCTCT ATGTGGAGAATGAGGGTGGGATGGGGTGCCCTTCATCCTGCGCTGCGCAAGGCCCTGAACGAGCGCAAGGCCG AGGTGAGGCTGCAGTTCCATGATGTGGCCGGCGACATCTTCCACCAGCAGTGCAAGCGCAACGAGCTGGTGATCC GCGTGCAGCCCAACGAGGCCGTGTACACCAAGATGATGACCAAGAAGCCGGGCATGTTCTTCAACCCCGAGGAGT CGGAGCTGGACCTACGGCAACAGATACAAGAACGTGAAGCTCCCTGACGCCTACGAGCGCCTCATCCTGG ACGTCTTCTGCGGGAGCCAGATGCACTTCGTGCGCAGCGACGAGCTCCGTGAGGCCTGGCGTATTTTCACCCCAC TGCTGCACCAGATTGAGCTGGAGAAGCCCAAGCCCATCCCCTATATTTATGGCAGCCGAGGCCCCACGGAGGCAG ACGAGCTGATGAAGAGAGGGGTTTCCAGTATGAGGGCACCTACAAGTGGGTGAACCCCCACAAGCTC**TGA**GCCC TGGGCACCCACCTCCACGCCACGGCCACCCTCCTTCCCGCCGCCCCGACCCCGAGTCGGGAGGACTCCGGGA GAGCCCAGCTACATTCCTCAGCTGCCAAGCACTCGAGACCATCCTGGCCCCTGCAGACCCTGAGCCCAGGA  $\verb|CCCACGTGAGAGAATCTGCCTGTGGCCTTGCCCGCCAGCCTCAGTGCCACTTGACATTCCTTGTCACCAGCAACA| \\$ TCTCGAGCCCCTGGATGTCCCCTGTCCCACCAACTCTGCACTCCATGGCCACCCCGTGCCACCCGTAGGCAGCC TCTCTGCTATAAGAAAAGCAGACGCAGCAGCTGGGACCCCTCCCAACCTCAATGCCCTGCCATTAAATCCGCAAA CAGCCAAAAAAAAAAAAAAAAAA

## 533/2825 FIGURE 480

MAEQVALSRTQVCGILREELFQGDAFHQSDTHIFIIMGASGDLAKKKIYPTIWWLFRDGLLPENTFIVGYARSRL
TVADIRKQSEPFFKATPEEKLKLEDFFARNSYVAGQYDDAASYQRLNSHMDALHLGSQANRLFYLALPPTVYEAV
TKNIHESCMSQIGWNRIIVEKPFGRDLQSSDRLSNHISSLFREDQIYRIDHYLGKEMVQNLMVLRFANRIFGPIW
NRDNIACVILTFKEPFGTEGRGGYFDEFGIIRDVMQNHLLQMLCLVAMEKPASTNSDDVRDEKVKVLKCISEVQA
NNVVLGQYVGNPDGEGEATKGYLDDPTVPRGSTTATFAAVVLYVENERWDGVPFILRCGKALNERKAEVRLQFHD
VAGDIFHQQCKRNELVIRVQPNEAVYTKMMTKKPGMFFNPEESELDLTYGNRYKNVKLPDAYERLILDVFCGSQM
HFVRSDELREAWRIFTPLLHQIELEKPKPIPYIYGSRGPTEADELMKRVGFQYEGTYKWVNPHKL

#### 534/2825 FIGURE 481

AGCTGAGGTGTGAGCAGCTGCCGAAGTCAGTTCCTTGTGGAGCCGGAGCTGGGCGCGGATTCGCCGAGGCACCGA GGCACTCAGAGGAGGTGAGAGAGCGGCGGCAGACAACAGGGGGACCCCGGGCCGGCCCAGAGCCGAGCCAAGC GTGCCCGCGTGTGTCCCTGCGTGTCCGCGAGGATGCGTGTTCGCGGGGTGTGTGCTGCGTTCACAGGTGTTTCTGC GGCAGGCGCC**ATG**TCAGAACCGGCTGGGGATGTCCGTCAGAACCCATGCGGCAGCAAGGCCTGCCGCCGCCTCTT CGGCCCAGTGGACAGCGAGCAGCTGAGCCGCGACTGTGATGCGCTAATGGCGGGCTGCATCCAGGAGGCCCGTGA GCGATGGAACTTCGACTTTGTCACCGAGACACCACTGGAGGGTGACTTCGCCTGGGAGCGTGTGCGGGGCCTTGG CTCACCTGCTCTGCTGCAGGGGACAGCAGAGGAAGACCATGTGGACCTGTCACTGTTGTACCCTTGTGCCTCG CTCAGGGGAGCAGGCTGAAGGGTCCCCAGGTGGACCTGGAGACTCTCAGGGTCGAAAACGGCGGCAGACCAGCAT  ${\tt GACAGATTTCTACCACTCCAAACGCCGGCTGATCTTCTCCAAGAGGAAGCCC} {\color{blue}{\textbf{TAA}}} {\tt TCCGCCCACAGGAAGCCTGC}$ AGTCCTGGAAGCGCGAGGGCCTCAAAGGCCCGCTCTACATCTTCTGCCTTAGTCTCAGTTTGTGTCTTAATTA TTATTTGTGTTTTAATTTAAACACCTCCTCATGTACATACCCTGGCCGCCCCCTGCCCCCAGCCTCTGGCATTA TACTTCCTCCTCCCACTTGTCCGCTGGGTGGTACCCTCTGGAGGGGTGTGGCTCCTTCCCATCGCTGTCACAGG CGGTTATGAAATTCACCCCCTTTCCTGGACACTCAGACCTGAATTCTTTTTCATTTGAGAAGTAAACAGATGGCA CTTTGAAGGGGCCTCACCGAGTGGGGGCATCATCAAAAACTTTGGAGTCCCCTCACCTCCTCTAAGGTTGGGCAG GGTGACCCTGAAGTGAGCACAGCCTAGGGCTGAGCTGGGGACCTGGTACCCTCCTGGCTCTTGATACCCCCCTCT GTCTTGTGAAGGCAGGGGGAAGGTGGGGTCCTGGAGCAGACCACCCCGCCTGCCCTCATGGCCCCTCTGACCTGC ACTGGGGAGCCCGTCTCAGTGTTGAGCCTTTTCCCTCTTTGGCTCCCCTGTACCTTTTGAGGAGCCCCAGCTACC  $\verb|CTTCTTCTCCAGCTGGGCTCTGCAATTCCCCTCTGCTGCTGTCCCTTGTCCCTTTCCCTTCAGTACCCTC| \\$  ${\tt TCAGCTCCAGGTGGCTCTGAGGTGCCTGTCCCACCCCCACCCCCAGCTCAATGGACTGGAAGGGGAAGGGACACA}$  ${\tt CAAGAAGAAGGGCACCCTAGTTCTACCTCAGGCAGCTCAAGCAGCCGACCGCCCCTCCTCTAGCTGTGGGGGTGA}$  ${\tt GGGTCCCATGTGGGGCACAGGCCCCCTTGAGTGGGGTTATCTCTGTGTTAGGGGTATATGATGGGGGAGTAGAT}$ CTTTCTAGGAGGAGACACTGGCCCCTCAAATCGTCCAGCGACCTTCCTCATCCACCCCATCCCCAGTTCA TTGCACTTTGATTAGCAGCGGAACAAGGAGTCAGACATTTTAAGATGGTGGCAGTAGAGGCTATGGACAGGGCAT GCCACGTGGGCTCATATGGGGCTGGGAGTAGTTGTCTTTCCTGGCACTAACGTTGAGCCCCTGGAGGCACTGAAG TGCTTAGTGTACTTGGAGTATTGGGGTCTGACCCCAAACACCTTCCAGCTCCTGTAACATACTGGCCTGGACTGT TTTCTCTCGGCTCCCCATGTGTCCTGGTTCCCGTTTCTCCACCTAGACTGTAAACCTCTCGAGGGCAGGGACCAC ACCCTGTACTGTTCTGTGTCTTTCACAGCTCCTCCCACAATGCTGAATATACAGCAGGTGCTCAATAAATGATTC TTAGTGACTTTAAAAAAAAAAAAAAAAAAAA

# 535/2825 FIGURE 482

 ${\tt MSEPAGDVRQNPCGSKACRRLFGPVDSEQLSRDCDALMAGCIQEARERWNFDFVTETPLEGDFAWERVRGLGLPKLYLPTGPRRGRDELGGGRRPGTSPALLQGTAEEDHVDLSLSCTLVPRSGEQAEGSPGGPGDSQGRKRRQTSMTDFYHSKRRLIFSKRKP$ 

#### 536/2825 FIGURE 483

 $\tt GTGGAACCTCGATATTGGTGGTGTCCATCGTGGGCAGCGGACTAATAAAGGCC\textbf{ATG}GCGCCAGCAGAAATCCTGA$ ACGGGAAGGAGTCTCCGCGCAAATAAGGGCGAGACTGAAAAATCAAGTCACTCAGTTGAAGGAGCAAGTACCTG GTTTCACACCACGCCTGGCAATATTACAGGTTGGCAACAGAGATGATTCCAATCTTTATATAAATGTGAAGCTGA AGGCTGCTGAAGAGATTGGGATCAAAGCCACTCACATTAAGTTACCAAGAACCACCAGAATCTGAGGTGATGA AGTACATTACATCTTTGAATGAAGACTCTACTGTACATGGGTTCTTAGTGCAGCTACCTTTAGATTCAGAGAATT GGAGACTTGCTAGAGGTGACCTCAATGACTGTTTCATTCCTTGTACGCCTAAGGGATGCTTGGAACTCATCAAAG AGACAGGGGTGCCGATTGCCGGAAGGCATGCTGTGGTGGTTGGGCCCCAGTAAAATAGTTGGGGCCCCGATGCATG ACTTGCTTCTGTGGAACAATGCCACAGTGACCACCTGCCACTCCAAGACTGCCCATCTGGATGAGGAGGTAAATA AAGGTGACATCCTGGTGGTTGCAACTGGTCAGCCTGAAATGGTTAAAGGGGGAGTGGATCAAACCTGGGGCAATAG TCATCGACTGTGGAATCAATTATGTCCCAGATGATAAAAAACCAAATGGGAGAAAAGTTGTGGGTGATGTGGCAT ACGACGAGGCCAAAGAGAGGGCGAGCTTCATCACTCCTGTTCCTGGCGGCGTAGGGCCCATGACAGTTGCAATGC TCATGCAGAGCACAGTAGAGAGTGCCAAGCGTTTCCTGGAGAAATTTAAGCCAGGAAAGTGGATGATTCAGTATA ACAACCTTAACCTCAAGACACCTGTTCCAAGTGACATTGATATATCACGATCTTGTAAACCGAAGCCCATTGGTA AGCTGGCTCGAGAAATTGGTCTGCTGTCTGAAGAGGTAGAATTATATGGTGAAACAAAGGCCAAAGTTCTGCTGT CAGCACTAGAACGCCTGAAGCACCGGCCTGATGGGAAATACGTGGTGGTGACTGGAATAACTCCAACACCCCTGG GAGAAGGGAAAAGCACAACTACAATCGGGCTAGTGCAAGCCCTTGGTGCCCATCTCTACCAGAATGTCTTTGCGT GTGTGCGACAGCCTTCTCAGGGCCCCACCTTTGGAATAAAAGGTGGCGCTGCAGGAGGCGGCTACTCCCAGGTCA TTCCTATGGAAGAGTTTAATCTCCACCTCACAGGTGACATCCATGCCATCACTGCAGCTAATAACCTCGTTGCTG TAAATGGAGTGAGAAGGTTCTCTGACATCCAAATCCGAAGGTTAAAGAGACTAGGCATTGAAAAGACTGACCCTA CCACACTGACAGATGAAGAGATAAACAGATTTGCAAGATTGGACATTGATCCAGAAACCATAACTTGGCAAAGAG TGTTGGATACCAATGATAGATTCCTGAGGAAGATCACGATTGGACAGGCTCCAACGGAGAAGGGTCACACACGGA CGGCCCAGTTTGATATCTCTGTGGCCAGTGAAATTATGGCTGTCCTGGCTCTCACCACTTCTCTAGAAGACATGA GAGAGAGACTGGGCAAAATGGTGGTGGCATCCAGTAAGAAAGGAGACCCGTCAGTGCCGAAGATCTGGGGGTGA GTGGTGCACTGACAGTGCTTATGAAGGACGCAATCAAGCCCAATCTCATGCAGACACTGGAGGGCACTCCAGTGT TTGTCCATGCTGGCCCGTTTGCCAACATCGCACATGGCAATTCCTCCATCATTGCAGACCAGATCGCACTCAAGC  $\tt TTGTTGGCCCAGAAGGGTTTGTAGTGACGGAAGCAGGATTTGGAGCAGACATTGGAATGGAAAAGTTTTTTAACA$ TCAAATGCCGGTATTCCGGCCTCTGCCCCCACGTGGTGGTGCTTGTTGCCACTGTCAGGGCTCTCAAGATGCACG AAGGCTTCAGTAACTTGAAGAAACAAATTGAAAATGCCAGAATGTTTGGAATTCCAGTAGTAGTGGCCGTGAATG CATTCAAGACGGATACAGAGTCTGAGCTGGACCTCATCAGCCGCCTTTCCAGAGAACATGGGGCTTTTGATGCCG TGAAGTGCACTCACTGGGCAGAAGGGGGCAAGGGTGCCTTAGCCCTGGCTCAGGCCGTCCAGAGAGCAGCACAAG CACCCAGCAGCTTCCAGCTCCTTTATGACCTCAAGCTCCCAGTTGAGGATAAAATCAGGATCATTGCACAGAAGA TCTATGGAGCAGATGACATTGAATTACTTCCCGAAGCTCAACACAAAGCTGAAGTCTACACGAAGCAGGGCTTTG GGAATCTCCCCATCTGCATGGCTAAAACACACTTGTCTTTGTCTCACAACCCAGAGCAAAAAGGTGTCCCTACAG GCTTCATTCTGCCCATTCGCGACATCCGCGCCAGCGTTGGGGCTGGTTTTCTGTACCCCTTAGTAGGAACGATGA GCACAATGCCTGGACTCCCCACCCGGCCCTGTTTTTATGATATTGATTTTGGACCCTGAAACAGAACAGGTGAATG GATTATTCTAAACAGATCACCATCCATCTTCAAGAAGCTACTTTGAAAGTCTGGCCAGTGTCTATTCAGGCCCAC TGGGAGTTAGGAAGTATAAGTAAGCCAAGAGAAGTCAGCCCCTGCCCAGAAGATCTGAAACTAATAGTAGGAGTT GTGACGTTCCACAGAATAAAAGGAAACAAGTTTGC

#### 537/2825 FIGURE 484

MAPAEILNGKEISAQIRARLKNQVTQLKEQVPGFTPRLAILQVGNRDDSNLYINVKLKAAEEIGIKATHIKLPRT
TTESEVMKYITSLNEDSTVHGFLVQLPLDSENSINTEEVINAIAPEKDVDGLTSINAGRLARGDLNDCFIPCTPK
GCLELIKETGVPIAGRHAVVVGRSKIVGAPMHDLLLWNNATVTTCHSKTAHLDEEVNKGDILVVATGQPEMVKGE
WIKPGAIVIDCGINYVPDDKKPNGRKVVGDVAYDEAKERASFITPVPGGVGPMTVAMLMQSTVESAKRFLEKFKP
GKWMIQYNNLNLKTPVPSDIDISRSCKPKPIGKLAREIGLLSEEVELYGETKAKVLLSALERLKHRPDGKYVVVT
GITPTPLGEGKSTTTIGLVQALGAHLYQNVFACVRQPSQGPTFGIKGGAAGGGYSQVIPMEEFNLHLTGDIHAIT
AANNLVAAAIDARIFHELTQTDKALFNRLVPSVNGVRRFSDIQIRRLKRLGIEKTDPTTLTDEEINRFARLDIDP
ETITWQRVLDTNDRFLRKITIGQAPTEKGHTRTAQFDISVASEIMAVLALTTSLEDMRERLGKMVVASSKKGEPV
SAEDLGVSGALTVLMKDAIKPNLMQTLEGTPVFVHAGPFANIAHGNSSIIADQIALKLVGPEGFVVTEAGFGADI
GMEKFFNIKCRYSGLCPHVVVLVATVRALKMHGGGPTVTAGLPLPKAYIQENLELVEKGFSNLKKQIENARMFGI
PVVVAVNAFKTDTESELDLISRLSREHGAFDAVKCTHWAEGGKGALALAQAVQRAAQAPSSFQLLYDLKLPVEDK
IRIIAQKIYGADDIELLPEAQHKAEVYTKQGFGNLPICMAKTHLSLSHNPEQKGVPTGFILPIRDIRASVGAGFL
YPLVGTMSTMPGLPTRPCFYDIDLDPETEQVNGLF

#### 538/2825 FIGURE 485

CGCGATTCTCAGGGATTGATCCGCCTCTTCAGGTAAGTTATCTTCCGGCCCCGTACCACTGTGCCACAGGCGCAG  $\verb|CCCGCTTCCTCAGGTGCCCTATCCCGCGCAGAAGACCACGGCTTCACAGAGTGTTATTTAAGGGCCGTGGCCAGCG|\\$ GAACATCCCGCCCCATTCTGTGACGCACGGGGTGGCCCCGTGGGACCCGAGGGGTGGGGCTGGGTTTAGTAGGA GACCIGGGGCAAGGCCCCCIGTGGACGACCATCTGCCAGCTTCTCTCGTTCCGTCGATTGGGAGGAGCGGTGGCG  ${ t ACCTCGGCCTTCAGTGTTTCCGACGGAGTGA} { t ATC} { t GCGGCGGCGGCTGGGATGCTGCTGCTGCTGCAGGC}$ GGGTGGGTCGGTGCTGGCCAGGCGATGGAGAAGGTGACAGGCGGCAACCTCTTGTCCATGCTGATCGCCTG CGCCTTCACCCTCAGCCTGGTCTACCTGATCCGTCTGGCCGCCGGCCACCTGGTCCAGCTGCCCGCAGGGGTGAA AAGTCCTCCATACATTTCTCCCCAATTCCATTCCTTGGGCATGCCATAGCATTTGGGAAAAGTCCAATTGAATT GGGGAGTGATGCTGCACTGCTTTTTAATAGTAAAAATGAAGACCTGAATGCAGAAGATGTCTACAGTCGCCT GACAACACCTGTGTTTGGGAAGGGAGTTGCATACGATGTGCCTAATCCAGTTTTCTTGGAGCAGAAGAAAATGTT AAAAAGTGGCCTTAACATAGCCCACTTTAAACAGCATGTTTCTATAATTGAAAAAGAAACAAAGGAATACTTTGA GAGTTGGGGGAGAAAGTGGGAGAAAAAATGTGTTTTGAAGCTCTTTCTGAGCTCATAATTTTAACAGCTAGCCATTG AATCAAGGATATTTTCTATAAGGCAATCCAGAAACGCAGACAGTCTCAAGAAAAATTGATGACATTCTCCAAAC TTTACTAGATGCTACATACAAGGATGGGCGTCCTTTGACTGATGATGAAGTAGCAGGGATGCTTATTGGATTACT  $\tt CTTGGCAGGGCAGCATACATCCTCAACTACTAGTGCTTGGATGGGCTTCTTTTTGGCCAGAGACAAAACACTTCA$ AAAAAAATGTTATTTAGAACAGAAAACAGTCTGTGGAGAGAATCTGCCTCCTTTAACTTATGACCAGCTCAAGGA TCTAAATTTACTTGATCGCTGTATAAAAGAAACATTAAGACTTAGACCTCCTATAATGATCATGATGAGAATGGC CAGAACTCCTCAGACTGTGGCAGGGTATACCATTCCTCCAGGACATCAGGTGTGTTTTCTCCCACTGTCAATCA AAGACTTAAAGACTCATGGGTAGAACGCCTGGACTTTAATCCTGATCGCTACTTACAGGATAACCCAGCATCAGG GGAAAAGTTTGCCTATGTGCCATTTGGAGCTGGGCGTCATCGTTGTATTGGGGAAAATTTTGCCTATGTTCAAAT TAAGACAATTTGGTCCACTATGCTTCGTTTATATGAATTTGATCTCATTGATGGATACTTTCCCACTGTGAATTA  ${\tt TACAACTATGATTCACACCCCTGAGAACCCAGTTATCCGTTACAAACGAAGATCAAAA} {\tt TGA} {\tt AAAAAGGTTGCAAGG}$ AACGAATATATGTGATTATCACTGTAAGCCACAAAGGCATTCGAAGAGAATGAAGTGTACAAAACAACTCTTGTA GTTTACTGTTTTTTTAAGTGTGTAATTCTAAAAGCCAGTTTATGATTTAGGATTTTGTTAACTGAATGGTTCTAT CAAATATAATAGCATTTGACACATTTTCTAATAGTTATGATACTTATACATGTGCTTTCAGGAAGTTCCTTGGTG AAACAATTGTTGAGGGGGGGATCTAGGTAATTGGCAGATTCTAAATAATATAATTTCCAGATAGTAATTTTAAGAG TACTCATCGCTCTTGCCAAATAAGTTCAGGGTATTCAAATCTTGGACTAGTCCTGCAAGGTATAAAGAATAAAAA GTAGGGAATAATCGAACATCTAAAGCCTTGAATAAGAGAATACTAATTGTTTTGGTATGATGATACTCAGAAATG GAGATATTATAGGAAAAAGAAATCCTTTGGAATTTTAACTAAAATCACTGCATATGGGAAATTAAGAGATCCAGG ACCATATTTGATAAGAGTTCCTAAAAATAATGTAATTATTAATGCTAAAGACTGCTCATGTATCTTGATCTAATT ACTAAATAAATTACATATTTATTTACCTGATAAATATGTATCTAGTTCTACAAGGTCACATTTATGTGGAAGTCC TTGAGATGGAGTCTCGCTCTGTTGCCCAGGCTGGAGTGCAGTGCTGCATCTCAGCTCACTGCATCCTCTGCCTC CCGGGTTCAAGCAATTCTCTGCCTCAGCCTCCCAAGTAGTTGGGATTACAGGCACCTGCCACCATGCCTGGCTAA TTTTTTGTATTTTAGTAGAGACGGGGGTTTCACTATGTTGGCTAGGCTGGTCTTGAACTCCTGAGCCTCGTGAG TCCACCCGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCCGCACCTGGCCTTCCCTGCTCCTCT GTGAGATCAGTGAACTCTGGTTTTAAGATAATCTGAAACAAGGTCCTTGGGAGTAATAAAATTGGTCACATTCTG TAAAGCACATTCTGTTTAGGAATCAACTTATCTCAAATTGTAACTCGGGGCCTAACTATATGAGATGGCTGAAAA AATACCACATCGTCTGTTTTCACTAGGTGATGCCAAAATATTTTGCTTTATGTATATTACAGTTCTTTTTAAAAC ACTGGAAGACTCATGTTAAACTCTAATTGTGAAGGCAGAATCTCTGCTAATTTTTCAGATTAAAATTCTCTTTGA AAAAAT

## 539/2825 FIGURE 486

MAAAAGMLLLGLLQAGGSVLGQAMEKVTGGNLLSMLLIACAFTLSLVYLIRLAAGHLVQLPAGVKSPPYIFSPIP FLGHAIAFGKSPIEFLENAYEKYGPVFSFTMVGKTFTYLLGSDAAALLFNSKNEDLNAEDVYSRLTTPVFGKGVA YDVPNPVFLEQKKMLKSGLNIAHFKQHVSIIEKETKEYFESWGESGEKNVFEALSELIILTASHCLHGKEIRSQL NEKVAQLYADLDGGFSHAAWLLPGWLPLPSFRRDRAHREIKDIFYKAIQKRRQSQEKIDDILQTLLDATYKDGR PLTDDEVAGMLIGLLLAGQHTSSTTSAWMGFFLARDKTLQKKCYLEQKTVCGENLPPLTYDQLKDLNLLDRCIKE TLRLRPPIMIMMRMARTPQTVAGYTIPPGHQVCVSPTVNQRLKDSWVERLDFNPDRYLQDNPASGEKFAYVPFGA GRHRCIGENFAYVQIKTIWSTMLRLYEFDLIDGYFPTVNYTTMIHTPENPVIRYKRRSK

## 540/2825 FIGURE 487

# 541/2825 FIGURE 488

 ${\tt MATVQQLEGRWRLVDSKGFDEYMKELGVGIALRKMGAMAKPDCIITCDGKNLTIKTESTLKTTQFSCTLGEKFEE} \\ {\tt TTADGRKTQTVCNFTDGALVQHQEWDGKESTITRKLKDGKLVVECVMNNVTCTRIYEKVE} \\$ 

#### 542/2825 FIGURE 489

#### 543/2825 FIGURE 490

 $\tt TGGCCGAAGCAGGGGACAGCAAGGGACGCTCAGGCGGGGACC\underline{ATG} GCGGACGGCGGCTCGGAGCGGCTGACGG$ GCGCATCGTCAAGATGGAGGTGGACTACAGCGCCACGGTGGATCAGCCCAAGTGTGCGAAGCTAGCCAA ATCGACATCCCGTATCTTAGTTGCAGTAGTGAAGATGTGCTATGAGGCTAAAGAATGGGATTTACTTAATGAAAA TATTATGCTTTTGTCCAAAAGGCGGAGTCAGTTAAAACAAGCTGTTGCCAAAATGGTTCAACAGTGCTGTACTTA TGTTGAGGAAATCACAGACCTTCCTATCAAACTTCGATTAATTGATACTCTACGAATGGTTACCGAAGGCAAGAT TTATGTTGAAATTGAGCGTGCGCGACTGACTAAAACATTAGCAACTATAAAAGAACAAAATGGTGATGTGAAAGA GGCAGCCTCCATTTTACAGGAGTTACAGGTGGAAACCTACGGGTCAATGGAAAAGAAGAAGAGCGAGTGGAATTTAT TTTGGAGCAAATGAGGCTCTGCCTAGCTGTGAAGGATTACATTCGAACACAAATCATCAGCAAGAAAATTAACAC CAAATTTTTCCAGGAAGAAATACAGAGAAATTAAAGTTGAAGTACTATAATTTAATGATTCAGCTGGATCAACA TGAGGGATCCTATTTGTCTATTTGTAAGCACTACAGAGCAATATATGATACTCCCTGTATACAGGCAGAAAGTGA AAAATGGCAGCAGGCTCTGAAGAGTGTTGTACTCTATGTTATCCTGGCTCCTTTTGACAATGAACAGTCAGATTT GGTTCACCGAATAAGTGGTGACAAGAAGTTAGAAGAAATTCCCAAATACAAGGATCTTTTAAAGCTTTTTACCAC AATGGAGTTGATGCGTTGGTCCACACTTGTTGAGGACTATGGAATGGAATTAAGAAAAGGTTCCCTTGAGAGTCC  ${\tt TGCAACGGATGTTTTTGGTTCTACAGAGGAAGGTGAAAAAAGGTGGAAAGACTTGAAGAACAGAGTTGTTGAACA}$ TGATGAGTCCGAAGCCTTTCTCTCAAATCTAGTAGTTAACAAGACCATCTTTGCTAAAGTAGACAGATTAGCAGG AATTATCAACTTCCAGAGACCCAAGGATCCAAATAATTTATTAAATGACTGGTCTCAGAAACTGAACTCATTAAT GTCTCTGGTTAACAAAACTACGCATCTCATAGCCAAAGAGGGGGTGATACATAATCTACAA**TAA**GGGTCTTAGTG  ${\tt TAAGCTTCTTTGTCTTAAATTTTAAAATAGTGAATATGTTTGAGACTCCCTTTGACCTTTCAGTTCCCCAAGTTCCCCAAGTTCCCCCCAAGTTCCCCCAAGTTCCCCCCAAGTTCCCCCAAGTTCCAAGTTCCAAGTTCCCCCAAGTTCCCCCAAGTTCCCCCCAAGTTCCCCCCCAAGTTCCCCCCAAGTTCCCCCAAG$ ATTGTTAACTTTGCATTTGCAATTGGTGCAAAAATACAGATTTCTGTCGTCTGAATACACAAAAAGTTGTGTCAT ATTCCAATAATAAAAGCAGTATATACATATTTCCTTTCTACAGTTACCTCTGATTCTCAACATTTTGTGGGGTAG TGATTTGGCAAGTGTTTTTTAAATAAAACAAATCTCATTGTAAAGTTATCAGTCATTTAGTAGAATAGAAAAGCA ACATAGAGCATACAAGAACATTTGGGATAGAGTTGTGATTTGTGAAGAATTTGTACTTTGATATTGTGGCGGAAA GTAGCTTGATTGCTTTCAGCCCTCATCCTCTCACTTGATCAGTTGTTCAACAGAATCAGCTGACATAATTGACAC CTCTCCCCTTTCTTTATATGGGTTTAAATTTAACATAAAGTTGTTTTTATAAGGCTTATTTGTGGCTTTAACTTG TAAGTCTGATTACATCATTATTGTTCCAAATTCATTATCTCTGTAGGAACTTTTAGTTCCATTATATGAACACTG GATAACCTAATTTTTTTAATGCTTTAAAAAAATGGCAAAAAGACGTCAGGCCACCCTCATAGTAAGTGGTGTAG TATTAAAATATTTCACGGAATTAAAAGTAGCTTGCTGTCAAAGAAACACCTGAGATGAATTGGTGTGAACGAAT TTTGCAAGTTTAATTTGATTTATTTCAGAGAAAATAGAAAAAACAATGTTAGAAGGTTATTTAAAATGATACTTA AATAAAGAAAGTGTGAGGTCTACTTTAAAAAAATTCAAATGAAGAAAAAAGAAAAACAGCATTCTAGAAATGGC ATTTCTCCTAATTAATTTCCACTTAATGGAAGATTATCAATTGTCCTATTTTATGATCCCAGGACTGAAGACAG TTGTGGGATATCTGTCATATTTATCCTGTGAGTCATTGTGAATAATGACATACAGTACTGAAGTAATCTGATTTT ATTCTTTGGAAATTCAATGCATTGGTCACACTAATAACATCAACATCTGCTATCACTTATCTTTTTAAAACTAAC TCTTTGTGTATTATGATTCAGTGCTATTTGTAACTGTGTCCCAGTGACCAAATTGCACTCGACTCGATCAGCTGT GTATTACAAAATAATTCATTAGCATAAAGTACAATAGTGAAATATTTGAGTTGTTCGGAACCTCAATTAATCCTG TTTTACATTTCAGACCTAAAGCTGGCAATCAGGAGAAGAAGCACTTTGTTTTAAATGTGGAGAAGATAACACTTG ATTCCATTTCATTGTCATTAGTGTATTAACCAGCAGGAGGGTGATGAGCCATTTTTCAAATGAAATACCTTTTA TTTCCATATAATTTTTTTTTTTTAGAGTTCAATAGCTGTTTCTATGATTATCCTCAATTTCCATATGTTACTGAA TCTGAAAACATCTTTAAAATTCAAACAGTTCCATTTTCTCTCTTGTAAGTGTTAAATGTGATAAAAGTACATAT TTTAAATTGTTTTCAGCTCTTGGATATAGCAGCAATAAAAACACTAATTTGTGGGTATTTAAGAAAACCTGGAGA ATAAACTCATACTTTAAAAGATC

# 544/2825 FIGURE 491

MADGGSERADGRIVKMEVDYSATVDQRLPECAKLAKEGRLQEVIETLLSLEKQTRTASDMVSTSRILVAVVKMCY EAKEWDLLNENIMLLSKRRSQLKQAVAKMVQQCCTYVEEITDLPIKLRLIDTLRMVTEGKIYVEIERARLTKTLA TIKEQNGDVKEAASILQELQVETYGSMEKKERVEFILEQMRLCLAVKDYIRTQIISKKINTKFFQEENTEKLKLK YYNLMIQLDQHEGSYLSICKHYRAIYDTPCIQAESEKWQQALKSVVLYVILAPFDNEQSDLVHRISGDKKLEEIP KYKDLLKLFTTMELMRWSTLVEDYGMELRKGSLESPATDVFGSTEEGEKRWKDLKNRVVEHNIRIMAKYYTRITM KRMAQLLDLSVDESEAFLSNLVVNKTIFAKVDRLAGIINFQRPKDPNNLLNDWSQKLNSLMSLVNKTTHLIAKEE MIHNLQ

#### 545/2825 FIGURE 492

TTGGTCTAAGACATGGGCTGACAAACTTTTTCCATAAAGGGCCAGAAAGTAAATATTTCAGGCTTTGTGGGATAT CTATTTATGATTTATTTTACTCCTAGTATTATTTCATCTTTTATACAGGTTATTAAGGTTAGTATTGCTGCATAA CAAATTACTCCAAAATTTAATCATTTATATTCAAGAAGAGTTCAGCTGCTAGGTTCTTATTTTGGGACTCTCATG GGGGCCTCCCAAATTCAAAAGGAAGAGACAAAGTCCACATCTGTCAATGCGAAGAATGGAAAGGTCACATATCAT ATGTGGAGCGGGACATAACAACTTCGGAAATTAAATCTCTTCCCCAGGTCAGTGTGCGGGCCTTCCACGCTGCCA GCGGAACACTGGAATGGCGGAAGGGGAACGGGTACGGCAGGGGGAAAGGCTGGGGGTGGGGTCGGGTCGAGTCGA GGGAAGCTGGCCCGAGTGTGGACAGAGATGGCGGTCTGCGCGTCTGTTGTTCCCAGCGCTCTGCGAAGCCTGAAA AGGAGGAGCAACCTGTCCAGAATCCCCGCAGGTCCGTGAAAGACAGGAAAAGGAGGGGGAAATCTCGACATGGAAA AACTCTACAGTGAAAATGAAGGAATGGCTTCAAACCAAGGAAAGATGGAAAATGAAGAACAGCCACAAGACGAGA GAAAGCCAGAAGTAACTTGTACTCTGGAAGACAAGAAGTTAGAAAACGAGGGGAAAGACAGAAAACAAGGGCAAAA CAGGAGATGAGGAAATGTTAAAGGATAAAGGAAAGCCAGAGAGTGAGGGAGAGGCAAAAGAAGGAAAGTCAGAGA AAGGAGACCAGGGAGTGAAACAAGGGCTGCAGGAAAGCCCCCAGCTGAGGATGATGTACCCAGGAAAGCCAAAA GAAAAACTAATAAGGGGCTGGCTCATTACCTCAAGGAGTATAAAGAGGCCATACATGATATGAATTTCAGCAATG AGGACATGATAAGAGAATTTGACAATATGGCTAAGGTGCAGGATGAGAAGAGAAAAAGCAAACAGAAATTGGGGG CGTTTTTGTGGATGCAAAGAAATTTACAGGACCCCTTCTACCCTAGAGGTCCAAGGGAATTCAGGGGTGGCTGCA TAACGTTACGGTAATACTTTACGTTAGGCATCCCTCCTGTTGCTAGCAGCCTTTTGACCTATCTGCAATGCAGTG TTCTCAGTAGGAAATGTTCATCTGTTACATGGAAAAAATGTTGATGGTGCATTGTAAAAATTAAAAAACACAACTT GCAGAACCAAAAAAAA

# 546/2825 **FIGURE 493**

MPAGGQAPQLTPVIPAFWEARGLPNSKGRDKVHICQCEEWKGHISYVERDITTSEIKSLPQVSVRAFHAASGTLE WRKGNGYGRGKGWGWGRVESREAGPSVDRDGGLRVCCSQRSAKPEKEEQPVQNPRRSVKDRKRRGNLDMEKLYSE NEGMASNQGKMENEEQPQDERKPEVTCTLEDKKLENEGKTENKGKTGDEEMLKDKGKPESEGEAKEGKSEREGES EMEGGSEREGKPEIEGKPESEGEPGSETRAAGKRPAEDDVPRKAKRKTNKGLAHYLKEYKEAIHDMNFSNEDMIR EFDNMAKVQDEKRKSKQKLGAFLWMQRNLQDPFYPRGPREFRGGCRAPRRDIEDIPYV

## 547/2825 FIGURE 494

ATAAATATTGGAAATGACAACTTTAACAATTCTATATGTACACAGGACACTGAAAACATAAAATCATGAACAAGG
CCAAAAAATAACGTTGCACATTAACCCTTTAGTTATTACTTTCTATTTTCCAGTCCCAGCATCATACCTGCTAAT
TACTCAGATCACAAGCCTCAAGGATTAAGTGTTTTGAATGTATTTCAGTTTCATACTTTAACAATGCTTAAAGAC
TATTGGTTGTATTCTGATCAAATGGTTCTCCCTTCCCATATTTCCCTTTCTCAATAGATCTTANACAAAAAGGTGA
GCAGGGAAAACTGTGATTACAGGATCCAACATGACAGGTATACCCAATGATTCATTTAAGGGTGGCATAAAGTTA
TATAGAAAAAGACCTACAGGACGGAAGACAAGTACATCAACATGTAGTGAAATTAAGCAGTTCACCCACTGCGTGC
TTACATGGCTCCTATATCTCTTCGGGAAGAAGCCAGAGCCGTAGAAAGTCAGGGCCAAACCAGAACTGTTGAGG
AGTATGGGTGGGGGGGCAGAAGGAAGGAAGGAAGGCAAAAAA

#### 548/2825 FIGURE 495A

GTACTCTCGCGGTATTTGTCCCGACTCTCGCGGGGTTTAGCGTGGCATTGGGAGGCCCGGCCTGGGGGAGGAC GGCGTTCCGTTAGCGGCGTTGGGGTTTGGCTGCAGTGCCTTTCTCTTCTGCTCACGGGGACCCGCTCAGG TCGAGATCAACCGGGAGCCGGTTGGTCGCATTATGTTTCAGCTCTTCTCAGACATATGTCCAAAAACATGCAAAA ACTTCCTTTGCTTGTGCTCAGGAGAGAAAGGCCTTGGGAAAACAACTGGGAAGAAGTTATGTTATAAAGGTTCTA CGTTCCATCGTGTGGTTAAAAACTTTATGATTCAGGGTGGGGACTTCAGTGAAGGTAATGGAAAAGGTGGAGAAT CAATTTATGGTGGATATTTTAAAGATGAAAACTTTATTCTCAAACATGACAGAGCGTTCCTTTTATCAATGGCAA ATCGAGGGAAACATACCAATGGTTCCCAGTTTTTCATTACCACAAAGCCTGCTCCACACCTGGATGGGGTGCATG TAGTCTTTGGACTGGTTATTTCTGGTTTTGAAGTAATCGAACAAATTGAAAATCTGAAGACCGATGCTGCAAGCA GACCATATGCAGATGTGCGAGTTATTGACTGTGGAGTACTTGCCACAAAATCAATAAAAGATGTTTTTGAGAAAA AAAGGAAGAAACCAACTCATTCAGAAGGCTCGGATTCCTCTTCCAATTCCTCCTCTTCTTCAGAATCATCTTCAG AAAGTGAACTTGAACATGAGAGAAGCAGAAGGAGGAAACATAAGAGGAGGCCAAAAGTTAAACGTTCTAAAAAGA GGCGAAAGGAAGCAAGCAGTTCAGAAGAGCCAAGGAATAAACATGCAATGAACCCAAAAGGTCACTCTGAGAGGA GTGATACCAATGAAAAAGGTCAGTTGATTCCAGTGCTAAAAGGGAAAAACCTGTGGTCCGCCCAGAAGAGATTC  $\verb|CTCCAGTGCCTGAGAACCGATTTTACTGAGAAGAGATATGCCTGTTGTTACTGCAGAACCTGAACCGAAGATTC| \\$ CAATTCGCTATCACACACCTCCAAGATCAAGATCCTGTTCTGAGTCAGATGATGATGACAGCAGTGAAACTCCTC CTCACTGGAAAGAGGAAATGCAGAGATTAAGAGCATATAGACCACCTAGTGGAGAAAAATGGAGTAAAGGAGATA AGTTAAGTGACCCCTGTTCAAGCCGATGGGATGAAAGAAGCTTGTCTCAGAGATCCAGATCATGGTCCTATAATG GATATTATTCAGACCTTAGTACAGCAAGACACTCTGGCCACCATAAAAAACGCAGAAAAGAAAAAAGGTTAAGC CTGACATAGAATCCTCAAAATCTTCCACTCGAAGAATGAAATCCTCTTGTGATAGAGAAAGGAGTTCTCGTTCTT CCTCATTGTCATCTCATCACTCATCAAAGAGAGACTGGTCTAAATCTGATAAGGATGTCCAGAGCTCTTTAACCC ATTCCAGCAGAGACTCATACAGATCAAAATCTCACTCACAGTCTTATTCTAGAGGAAGCTCAAGATCAAGGACTG CGTCAAAGTCCTCATCACATTCTCGAAGTAGATCAAAGTCCAGATCTAGTTCCAAGTCTGGGCACCGAAAGAGAG CATCAAAATCACCAAGAAAAACAGCTTCTCAGTTAAGTGAAAATAAACCAGTTAAAACAGAACCTTTAAGAGCAA CCATGGCACAAAATGAAAATGTAGTAGTACAACCAGTTGTAGCAGAAAATATTCCTGTAATACCACTGAGTGACA GTCCCCCCCTTCAAGATGGAAGCCTGGACAGAAACCTTGGAAGCCCTCTTATGAGCGAATTCAGGAAATGAAAG  $\tt CTAAAACAACCCATTTGCTACCCATCCAAAGCACTTACAGTTTAGCAAATATTAAAGAGACTGGTAGCTCATCAT$ CCTACCATAAAAGAGAAAAAATTCGGAAAGTGATCAGAGCACTTATTCAAAATACAGTGATAGAAGTTCAGAAA GCTCACCAAGGTCAAGGAGCAGATCTTCTAGGAGTAGATCTTATTCCAGATCATATACAAGATCACGTAGTCTAG GTAGTAGATCATCTTCATATACTTCTATTAGCAGTGATGATGGAAGGCGAGCTAAGAGGAGACTTAGATCCAGTG GGAAAAAAAATAGCGTTTCACATAAAAAGCATAGCAGCAGCTCTGAAAAGACACTTCACAGTAAATATGTCAAAG GTAGAGACAGGTCTTCATGTGTGAGAAAGTATAGCGAGAGCAGATCATCTTTAGATTATTCTTCAGACAGTGAGC AGTCAAGTGTTCAGGCCACAGTCAGCCCAGGAAAAAGAGAAGCAGGGCCAAATGGAAAGAACACATAATAAAC  ${ t A}{ t G}{ t A}{ t G}{ t C}{ t T}{ t C}{ t C}{ t C}{ t C}{ t A}{ t A}{ t A}{ t C}{ t C}{ t C}{ t A}{ t A}{ t A}{ t G}{ t C}{ t$ AGGCCACATCCGATTCTGAATCAGAGGTTAGTGAAATTCACATCAAAGTCAAACCCACAACCAAGTCGTCCACAA ATACTTCACTGCCTGATGATAATGGTGCTTGGAAATCAAGCAAACAGCGCACATCAACTTCTGACTCTGAGGGGT CCTGTTCCAATTCGGAAAACAATAGGGGAAAGCCACAAAAGCACAAACATGGGTCAAAGGAAAATCTTAAAAGAG AACACACCAAAAAGTGAAAGAGAAATTGAAAGGGAAAAAAGACAAAAAGCATAAGGCTCCAAAACGAAAGCAAG AATCAAAAGAGAAAAAGTTTCTGAAAAACAATGAAACCATAAAAGATAATATTCTAAAAACTGAGAAATCCAGTG AAGAGGACCTTTCAGGTAAACATGATACAGTGACTGTTTCATCAGATCTTGATCAGTTTACTAAAGATGATAGTA AACTCAGTATTTCTCCCACAGCTTTAAATACTGAGGAAAATGTGGCCTGTTTACAAAACATTCAGCACGTTGAAG AAAGTGTTCCCAATGGAGTGGAAGATGTGCTTCAAACAGATGACAACATGGAGATCTGCACTCCTGATAGGAGTT

#### 549/2825 FIGURE 495B

CCCCAGCAAAAGTAGAGGAGACTTCCCCTCTAGGAAATGCACGGCTTGATACCCCAGATATAAACATTGTTTTGA AGCAGGATATGGCAACGGAACATCCTCAAGCAGAGGTAGTAAAACAGGAAAGCAGCATGTCCGAAAGTAAAGTGT TGGGTGAAGTGGGGAAACAGGACAGCAGCTCTGCTAGCTTGGCTAGTGCTGGAGAAAGTACCGGGAAGAAGGAGG TGGCTGAGAAGAGCCAGATCAACCTCATTGATAAGAAATGGAAGCCCCTGCAAGGTGTGGGGAACCTGGCAGCAC CTAATGCTGCCACATCCAGTGCTGTGGAAGTTAAGGTGTTGACCACTGTGCCTGAAATGAAACCACAAGGCTTGA ACCGTAGACCAAGAAATCAGGAGAGTTCAAGTGATGAGCAGACGCCTAGTCGGGATGATGATAGCCAGTCCAGGA GAAGTCGATCGAGAAGTTCCACATCATCTTATCGATCAAGAAGCTACTCTAGAAGTCGGAGCAGAGGATGGTACA GCAGAGGCCGAACCAGAAGCCGGAGCAGTTCCTACCGGAGTTACAAAAGTCACAGGACGTCCAGCAGGAGCAGAT CCAGGAGCAGCTCATATGATCCCCACAGTCGATCCAGGTCCTACACCTACGATAGCTACTATAGCAGGAGTCGGA GTCGAAGTAGAAGCCAGAGAGTGACAGTTACCACCGAGGCAGAAGTTATAATCGGCGGTCCAGGAGTTGTAGAT  $\tt CTTATGGCTCTGACAGTGAAAGTGACCGAAGTTACTCTCATCACCGGAGCCCCAGTGAGAGCAGCAGATACAGT{\bf T}$  $\underline{\textbf{GA}} \texttt{AAACGTCCGGATACAAATTATATCTTATTTGTAAATATCTGGCAACTTAGCTTAAGAAATGTAATGACAGTCT}$ GTTGTTCTATTTCAATATCAGAGGTGAATTTCAAAAATAGACACTTCTTAATTGTTACTGGTTCATTTACATGTG GGGAGAAGAATTTAAAATACAGATATGTCTCCTAAAAATATTTTTTATGCCACATTTTACAGTAGCCAACTATGGA AATGAATTTCATTTTCTTGAATCAAGAAATCGTGAAATTTATCTATGTATAATTTGCAATATTATTTTAAGTCTA  $\tt TTTCACTCTATCTTACGTATCCCTTAGAATACAGATTCTTTTTGCCTGTTTTTCCAGTTTTAGCATATATGCTGC$ CAAGCATAGAACTGTGAAGGAGAACTGTTAAAGGCGGCCAAATATTTATATACTGATTACATAGAGTCTTGTACA TATGTGCTCTAAAAACAAACCACCCAGAATTGATACTGTTGGTAACCAGGAGTATAAGGCAGTGGCTCTGGGGTT CTTAATTCATTCCTAACTTCTTTGATACTTCACAGGATTAGGAAAGTGGTCATCATACATCCCACACAGTCTGTA CTTACTATTTTATGTTCTTTTTAATCAATACCGATCAGAAGTTTAGGTTATAAAAACAATTCTACTTCATGCTTT GGTGCTTGGTAATTTTTGGTGCGTCTTTAAGCATTACTCTTATATATCATATATTAAAAATACCATAAAAATGAAA TTCAGACAAAATCACTGGCACCAAAAATGGTTTATTCTGAGCTGTCTTCACTTTGACTATTTGGGGGGGCTTCTCT TGTGTGTGTGTGCCTCTGGAAGAGTGAAGAATGGACTTCAAAAGTAACATCAAAAATCTAACTGCCACCATCCTG GAGACATTTTGCAGGGCTTTCCTTTCAAGTCTTTCAAGTACAGGATATTACCACAACAGCAGCTGAACTGTTGTA AGACAGGGGAAGAGGGATAGAGGGTATAGCATAAATTACATATTTTCATGGCTTTGGGTGGTTCCTCCAAAAATA TTTGAATTTGTTCCCCAAGTGTACTTAATCACCTTAGTGCCAGTTTAATCCAGTTATGCAGAAGAAATTCATATT GGTTGCCTGATGTAGAGCTCAGCACCACCCTACCACAGGCCTTGTCTGGTGTATTTGGGAAGTGGAAAAGAGCCC GTATGAAAACAGGGTAAGGTGGTCAGTTGTTTGCCAGGTCAATAGACAGAAAGTACATTAGAAAACAGGACTTAG GCCAAACAAACAATACTGGATACTGAATACAAAACAGTATGATTTATATTAAAGGTTTCCAAAGGTTGCCTGCAA AGGAGAATATTACTACTAGTCAGCAGGAAAAAATGCATTCAGAACCCAAGCAGAAACTGCCAAATGTAATTAGG TTAAGAAAAGTTACCCTTGGGCAGTGTATTAGTTTTCTATTGCTGTGTGACAAATTACCCCAAATTTAGCAGTTT AAAAAACAATACCCATAGCAGTTCTGTAGCTCATGAGTCTGGCACAGTGTGGCTGGATTCTCTGCTCAGGGTCTT AAAGGCTGAAATAAGGGTTGGCAGGACAACATTCCTTCATGGAGGCTCTGGGGAAGAATCTGCTTCTAAGTTCAT CCTCCTGTTTTAGGGGCATATGATTAGCTCAAGCCCACAGATATATTTTAAGGTTGATTGTGCGATAGAACATAA TTGCAGGAGTACTGTCTCATCTCATCATATTCACGGGTTCTGGAGATTAGCTCATTGAAAGTGGGAGGGGCATTT  ${\tt TCAAATTCTGCCTACCACAGGCAATAACTGCCCATCTCAGCTGTAGGTGGAATTTTTACCCAGAAAAGATAGGCC}$ CTAGAAGCCTCATTTCTTCTCCATGGAAAAGGACAGCCCTCTGCTGCAGCGTTCAACTTGTGTTTACTGAC AGAGTGAACTACAGAAATAGCTTTTCTTCCTAAAGGGGGATTGTTCTACATTTTGAAGTTATTTTTTAATAAAATT GAATTATGTTGT

#### 550/2825 FIGURE 496

MGAQDRPQCHFDIEINREPVGRIMFQLFSDICPKTCKNFLCLCSGEKGLGKTTGKKLCYKGSTFHRVVKNFMIQG GDFSEGNGKGGESIYGGYFKDENFILKHDRAFLLSMANRGKHTNGSQFFITTKPAPHLDGVHVVFGLVISGFEVI EQIENLKTDAASRPYADVRVIDCGVLATKSIKDVFEKKRKKPTHSEGSDSSSNSSSSSESSESELEHERSRRRK  ${\tt HKRRPKVKRSKKRRKEASSSEEPRNKHAMNPKGHSERSDTNEKRSVDSSAKREKPVVRPEEIPPVPENRFLLRRD}$ MPVVTAEPEPKIPDVAPIVSDQKPSVSKSGRKIKGRGTIRYHTPPRSRSCSESDDDDSSETPPHWKEEMORLRAY RPPSGEKWSKGDKLSDPCSSRWDERSLSQRSRSWSYNGYYSDLSTARHSGHHKKRRKEKKVKHKKKGKKOKHCRR HKQTKKRRILIPSDIESSKSSTRRMKSSCDRERSSRSSSLSSHHSSKRDWSKSDKDVQSSLTHSSRDSYRSKSHS QSYSRGSSRSRTASKSSSHSRSRSKSRSSSKSGHRKRASKSPRKTASQLSENKPVKTEPLRATMAQNENVVVQPV VAENIPVIPLSDSPPPSRWKPGQKPWKPSYERIQEMKAKTTHLLPIQSTYSLANIKETGSSSSYHKREKNSESDO STYSKYSDRSSESSPRSRSRSRSRSYSRSYTRSRSLASSHSRSRSPSSRSHSRNKYSDHSQCSRSSSYTSISSD DGRRAKRRLRSSGKKNSVSHKKHSSSSEKTLHSKYVKGRDRSSCVRKYSESRSSLDYSSDSEQSSVQATQSAQEK EKQGQMERTHNKQEKNRGEEKSKSERECPHSKKRTLKENLSDHLRNGSKPKRKNYAGSKWDSESNSERDVTKNSK NDSHPSSDKEEGEATSDSESEVSEIHIKVKPTTKSSTNTSLPDDNGAWKSSKQRTSTSDSEGSCSNSENNRGKPQ KHKHGSKENLKREHTKKVKEKLKGKKDKKHKAPKRKQAFHWQPPLEFGEEEEEIDDKQVTQESKEKKVSENNET IKDNILKTEKSSEEDLSGKHDTVTVSSDLDQFTKDDSKLSISPTALNTEENVACLQNIQHVEESVPNGVEDVLQT DDNMEICTPDRSSPAKVEETSPLGNARLDTPDINIVLKQDMATEHPQAEVVKQESSMSESKVLGEVGKQDSSSAS LASAGESTGKKEVAEKSQINLIDKKWKPLQGVGNLAAPNAATSSAVEVKVLTTVPEMKPQGLRIEIKSKNKVRPG SLFDEVRKTARLNRRPRNQESSSDEQTPSRDDDSQSRSPSRSRSKSETKSRHRTRSVSYSHSRSRSSTSSYRS RSYSRSRSGWYSRGRTRSRSSSYRSYKSHRTSSRSRSRSSSYDPHSRSRSYTYDSYYSRSRSRSRSQRSDSYHR GRSYNRRSRSCRSYGSDSESDRSYSHHRSPSESSRYS

#### 551/2825 FIGURE 497A

CGGCAGGGTTGGAAAATGATGGAAGAGGCGAGGTGGAGGCGACCGAGTGCTGAGAGGAACCTGCGGAATCGGCC AGATTAACTAGAGAAAGAAGAAGCCCCTAGGCCCTATTCAAAACAAGTATCTTATGTTATTCAGGCTGAAGGA AAAGAGCATATTATTCACTTGGAAAGGAACAAAGACCTTTTGCCTGAAGATTTTGTGGTTTATACTTACAACAAG AATTCATCCATTGCTCTTAGCGACTGTTTTGGACTCAGAGGATTGCTGCATTTAGAGAATGCGAGTTATGGGATT GAACCCCTGCAGAACAGCTCTCATTTTGAGCACATCATTTATCGAATGGATGATGTCTACAAAGAGCCTCTGAAA TGTGGAGTTTCCAACAAGGATATAGAGAAAGAAACTGCAAAGGATGAAGAGGAAGAGCCTCCCAGCATGACTCAG CTACTTCGAAGAAGAAGACCTGTCTTGCCACAGACCCGGTATGTGGAGCTGTTCATTGTCGTAGACAAGGAAAGG TATATTATGTTAAATATTCGAATTGTGCTAGTTGGACTGGAGATTTGGACCAATGGAAACCTGATCAACATAGTT GGGGGTGCTGGTGATGTGCTGGGGAACTTCGTGCAGTGGCGGAAAAGTTTCTTATCACACGTCGGAGACATGAC AGTGCACAGCTAGTTCTAAAGAAAGGTTTTGGTGGAACTGCAGGAATGGCATTTGTGGGAACAGTGTGTTCAAGG AGCCACGCAGGCGGGATTAATGTGTTTGGACAAATCACTGTGGAGACATTTGCTTCCATTGTTGCTCATGAATTG GGTCATAATCTTGGAATGAATCACGATGATGGGAGAGATTGTTCCTGTGGAGCAAAGAGCTGCATCATGAATTCA GGAAAAACCAGTGAGTGTGATGTTCCAGAGTACTGCAATGGTTCTTCTCAGTTCTGTCAGCCAGATGTTTTTATT CAGAATGGATATCCTTGCCAGAATAACAAAGCCTATTGCTACAACGGCATGTGCCAGTATTATGATGCTCAATGT CAAGTCATCTTTGGCTCAAAAGCCAAGGCTGCCCCCAAAGATTGTTTCATTGAAGTGAATTCTAAAGGTGACAGA TTTGGCAATTGTGGTTTCTCTGGCAATGAATACAAGAAGTGTGCCACTGGGAATGCTTTGTGTGGAAAGCTTCAG TGTGAGAATGTACAAGAGATACCTGTATTTGGAATTGTGCCTGCTATTATTCAAACGCCTAGTCGAGGCACCAAA TGTTGGGGTGTGGATTTCCAGCTAGGATCAGATGTTCCAGATCCTGGGATGGTTAACGAAGGCACAAAATGTGGT GCTGGAAAGATCTGTAGAAACTTCCAGTGTGTAGATGCTTCTGTTCTGAATTATGACTGTGATGTTCAGAAAAAG TGTCATGGACATGGGGTATGTAATAGCAATAAGAATTGTCACTGTGAAAATGGCTGGGCTCCCCCAAATTGTGAG AGAAGCTACTTCAGAAAGAAGATCACAAACATATGAGTCAGATGGCAAAAATCAAGCAAACCCTTCTAGACAG CCGGGGAGTGTTCCTCGACATGTTTCTCCAGTGACACCTCCCAGAGAAGTTCCTATATATGCAAACAGATTTGCA GTACCAACCTATGCAGCCAAGCAACCTCAGCAGTTCCCATCAAGGCCACCTCCACCACAACCGAAAGTATCATCT  ${\tt CAGGGAAACTTAATTCCTGCCCGTCCTGCTCCTGCACCTCCTTTATATAGTTCCCTCACT} {\color{red}{\bf TGA}} {\tt TTTTTTTAACCT}$ GTTGCAACTATGAATGAAAACAAAACACCACAAAACAGACTTCACTAACACAGAAAACAGAAACTGAGTGTGAG AGTTGTGAAATACAAGGAAATGCAGTAAAGCCAGGGAATTTACAATAACATTTCCGTTTCCATCATTGAATAAGT CTTATTCAGTCATCGGTGAGGTTAATGCACTAATCATGGATTTTTTGAACATGTTATTGCAGTGATTCTCAAATT AACTGTATTGGTGTAAGATTTTTGTCATTAAGTGTTTAAGTGTTATTCTGAATTTTCTACCTTAGTTATCATTAA TGTAGTTCCTCATTGAACATGTGATAATCTAATACCTGTGAAAACTGACTAATCAGCTGCCAATAATATCTAATA ATTGTCTTCAAAAGAATGCACAAGAACCACAATTAAGATGTCATATTATTTTTGAAAGTACAAAATATACTAAAAG AGTGTGTGTGTATTCACGCAGTTACTCGCTTCCATTTTTATGACCTTTCAACTATAGGTAATAACTCTTAGAGAA ATTAATTTAATATTAGAATTTCTATTATGAATCATGTGAAAGCATGACATTCGTTCACAATAGCACTATTTTAAA TAAATTATAAGCTTTAAGGTACGAAGTATTTAATAGATCTAATCAAATATGTTGATTCATGGCTATAATAAAGCA GGAGCAATTATAAAATCTTCAATCAATTGAACTTTTACAAAACCACTTGAGAATTTCATGAGCACTTTAAAATCT GAACTTTCAAAGCTTGCTATTAAATCATTTAGAATGTTTACATTTACTAAGGTGTGCTGGGTCATGTAAAATATT 

# 552/2825 FIGURE 497B

#### 553/2825 FIGURE 498

MGSGARFPSGTLRVRWLLLLGLVGPVLGAARPGFQQTSHLSSYEIITPWRLTRERREAPRPYSKQVSYVIQAEGK EHIIHLERNKDLLPEDFVVYTYNKEGTLITDHPNIQNHCHYRGYVEGVHNSSIALSDCFGLRGLLHLENASYGIE PLQNSSHFEHIIYRMDDVYKEPLKCGVSNKDIEKETAKDEEEEPPSMTQLLRRRRAVLPQTRYVELFIVVDKERY DMMGRNQTAVREEMILLANYLDSMYIMLNIRIVLVGLEIWTNGNLINIVGGAGDVLGNFVQWREKFLITRRHDS AQLVLKKGFGGTAGMAFVGTVCSRSHAGGINVFGQITVETFASIVAHELGHNLGMNHDDGRDCSCGAKSCIMNSG ASGSRNFSSCSAEDFEKLTLNKGGNCLLNIPKPDEAYSAPSCGNKLVDAGEECDCGTPKECELDPCCEGSTCKLK SFAECAYGDCCKDCRFLPGGTLCRGKTSECDVPEYCNGSSQFCQPDVFIQNGYPCQNNKAYCYNGMCQYYDAQCQ VIFGSKAKAAPKDCFIEVNSKGDRFGNCGFSGNEYKKCATGNALCGKLQCENVQEIPVFGIVPAIIQTPSRGTKC WGVDFQLGSDVPDPGMVNEGTKCGAGKICRNFQCVDASVLNYDCDVQKKCHGHGVCNSNKNCHCENGWAPPNCET KGYGGSVDSGPTYNEMNTALRDGLLVFFFLIVPLIVCAIFIFIKRDQLWRSYFRKKRSQTYESDGKNQANPSRQP GSVPRHVSPVTPPREVPIYANRFAVPTYAAKQPQQFPSRPPPPQPKVSSQGNLIPARPAPAPPLYSSLT

#### 554/2825 FIGURE 499

GAACCGTATCATCCAGTTTAACCCAGGGCCAGAGAAGTACTTCACCCTGGGGGCTCCCCACTGGGAGTACCCCACT TGGCTGCTACAAGAAGCTGATTGAATACTATAAGAATGGGGACCTGTCCTTTAAATATGTGAAGACCTTCAACAT GGATGAGTACGTGGGCCTTCCTCGAGACCACCCGGAGAGTTACCACTCCTTCATGTGGAACAACTTCTTCAAGCA CATTGACATCCACCCAGAAAACACCCACATTCTGGATGGGAATGCAGTCGACCTACAGGCAGAATGTGATGCCTT TGAAGAGAAGATCAAGGCTGCAGGTGGGATCGAGCTATTTGTTGGAGGCATCGGCCCTGATGGACACATTGCCTT CAACGAGCCAGGCTCCAGTCTGGTGTCCAGGACCCGTGTGAAGACGCTGGCCATGGATACCATCCTGGCCAATGC TAGGTTCTTCGATGGAGAACTCACCAAGGTGCCCACCATGGCCTTGACGGTGGGGGTGGGCACTGTCATGGATGC GCTGAAAGTGAAGACTGTCAAGTATTTCAAAGGTTTAATGCTTGTTCATAACAAGTTGGTGGACCCCTTGTACAG TATCAAAGAGAAAGAAACTGAGAAAAGCCAATCTTCGAAGAAACCATACAGCGAT**TAG**CCTGTGCTGGGACCTAG AGTATGGTTACTCCAGATAAGTGGGTGAACTTATTGTTCTTGGCCATGAGGCTGGGAGCCTAGTCACGGAGTTTA GCTATAGGGAGAATGTTTGTAACTTAATCAGAAAAAAATATCTGCAAAATGTACTCCATCATTTTGATGTCTGC CAAACCCAGGTTGGGAGTTTTAAACTTTTTGTTCTGCTTCAGCCATGGTTCACACATATGACACACTCCCGTCAG TGAGAGTCCTTCTTGCACAGGCCTGCCCCTTGGTTGAGAACCATTGTTCCAAGTGAAGGCACAAACTCTCAATAT CTATTGATAAATCCAGAGCCTCAAGGGGCCCAGCCACGTCAAACTTCTCTCCCTCAGGGACTCTCCAGCACCAAA AGGCAGAAGGTGGAAGCCGTTTTTCCCCCAGAGCCCTGTGTTTTTGTGAAAGGCCTCACTGTGGCTCCTCTGTTT TACATACTCATTAGTAAGTGGGAGGTCCACTGGGGCAACAGACACTGCCACAATTTCAGTGTTGTGTTCAGCCAA GGGGACGGTCTGGACAGCCTTAAGTGTGAGTTTAGTCACAACTCCTGAGTGTCCCGCTCTCCTGCTTACCTA GGAGGTGAGTGCCAGGAAAATACACCAAATGCTTCTAGTATTGTTTCCCCACTTAAAATAGTCCTGCTTAAATTC ACATGGTGTGTGTCTGAGGAGCATCAGGAAATACAACCCTTTTGCCCATTTACCCTTCTCCCCGGATCC  $\verb|TTTTGTGCCTGTTTGAAGCTACTGCTGCCTCCATTTCTGGGAAAGACCTTTGAGAGCCTAGCCCAGGCCTAAGG|$ GCTATGTTTGGTACCAGTGTTTTGTCTTTAGCTTTTCTATGTGATTGTGCCGTCATTCTGTTTTAAGCTCATGGA 

# 555/2825 FIGURE 500

MKLIILEHYSQASEWAAKYIRNRIIQFNPGPEKYFTLGLPTGSTPLGCYKKLIEYYKNGDLSFKYVKTFNMDEYV GLPRDHPESYHSFMWNNFFKHIDIHPENTHILDGNAVDLQAECDAFEEKIKAAGGIELFVGGIGPDGHIAFNEPG SSLVSRTRVKTLAMDTILANARFFDGELTKVPTMALTVGVGTVMDAREVMILITGAHKAFALYKAIEEGVNHMWT VSAFQQHPRTVFVCDEDATLELKVKTVKYFKGLMLVHNKLVDPLYSIKEKETEKSQSSKKPYSD

#### 556/2825 FIGURE **501**

AAAACAGCCGGGGCTCCAGCGGGAGAACGATA**ATG**CAAAGTGCTATGTTCTTGGCTGTTCAACACGACTGCAGAC CCATGGACAAGAGCGCAGGCAGTGGCCACAAGAGCGAGGAGAAAGCGAAAAGATGAAACGGACCCTTTTAAAAG ATTGGAAGACCCGTTTGAGCTACTTCTTACAAAATTCCTCTACTCCTGGGAAGCCCAAAAACCGGCAAAAAAGCA AACAGCAAGCTTTCATCAAGCCTTCTCCTGAGGAAGCACAGCTGTGGTCAGAAGCATTTGACGAGCTGCTAGCCA CCTGTGAAGACTTCAAAAAAACCAAATCACCCCAAAAGCTGTCCTCAAAAGCAAGGAAAATATATACTGACTTCA TAGAAAAGGAAGCTCCAAAAGAGATAAACATAGATTTTCAAACCAAAACTCTGATTGCCCAGAATATACAAGAAG CTACAAGTGGCTGCTTTACAACTGCCCAGAAAAGGGTATACAGCTTGATGGAGAACAACTCTTATCCTCGTTTCT TGGAGTCAGAATTCTACCAGGACTTGTGTAAAAAGCCACAAATCACCACAGAGCCTCATGCTACA**TGA**AATGTAA AAGGGAGCCCAGAAATGGAGGACATTTCATTCTTTTTCCTGAGGGGAAGGACTGTGACCTGCCATAAAGACTGAC CTTGAATTCAGCCTGGGTGTTCAGGAACATCACTCAGAACTATTGATTCAAAGTTGGGTAGTGAATCAGGAAGC CAGTAACTGACTAGGAGAAGCTGGTATCAGAACAGCTTCCCTCACTGTGTACAGAACGCAAGAAGGGGAATAGGTG GTCTGAACGTGGTGTCTCACTCTGAAAAGCAGGAATGTAAGATGATGAAAGAGACAATGTAATACTGTTGGTCCA AAAGCATTTAAAATCAATAGATCTGGGATTATGTGGCCTTAGGTAGCTGGTTGTACATCTTTCCCTAAATCGATC CATGTTACCACATAGTAGTTTTAGGTTTAGGATTCAGTAACAGTGAAGTGTTTACTATGTGCAAGGGTATTGAAGT TCTTATGACCACAGATCATCAGTACTGTTGTCTCATGTAATGCTAAAACTGAAATGGTCCGTGTTTGCATTGTTA AAAATGATGTGTGAAATAGAATGAGTGCTATGGTGTTGAAAACTGCAGTGTCCGTTATGAGTGCCAAAAATCTGT 

# 557/2825 FIGURE 502

MQSAMFLAVQHDCRPMDKSAGSGHKSEEKREKMKRTLLKDWKTRLSYFLQNSSTPGKPKTGKKSKQQAFIKPSPE EAQLWSEAFDELLASKYGLAAFRAFLKSEFCEENIEFWLACEDFKKTKSPQKLSSKARKIYTDFIEKEAPKEINI DFQTKTLIAQNIQEATSGCFTTAQKRVYSLMENNSYPRFLESEFYQDLCKKPQITTEPHAT

### 558/2825 FIGURE 503

GAGAAAAGTGATTTTGAAGCTGTAGAAGCACTTATGTCAATGAGCTGCAGTTGGAAGTCTGATTTTAAGAAATAC GTTGAAAACAGACCTGTTACACCAGTATCTGATTTGTCAGAGGAAGAGAATCTGCTTCCGGGAACACCTGATTTT CATACAATCCCAGCATTTTGTTTGACTCCACCTTACAGTCCTTCTGACTTTTGAACCCTCTCAAGTGTCAAATCTG ATGGCACCAGCGCCATCTACTGTACACTTCAAGTCACTCTCAGATACTGCCAAACCTCACATTGCCGCACCTTTC AAAGAGGAAGAAAAGAGCCCAGTATCTGCCCCCAAACTCCCCAAAGCTCAGGCAACAAGTGTGATTCGTCATACA TCCAAATGTGAGAAACACAGTGGCAGATGTTGATGAGAAAGCAAGTGCTGCACTTTATGACTTTTCTGTGCCT TCCTCAGAGACGGTCATCTGCAGGTCTCAGCCAGCCCCTGTGTCCCCACAACAGAAGTCAGTGTTGGTCTCTCCA CCTGCAGTATCTGCAGGGGGAGTGCCACCTATGCCGGTCATCTGCCAGATGGTTCCCCTTCCTGCCAACAACCCT ACACAAGTCCCCAAAGGCGCTGTCATGTTTGTGGTACCCCAGCCCGTTGTGCAGAGGTTCAAAGCCTCCGGTGGTG AGCCCGAATGGCACCAGACTCTCCCCATTGCCCCTGCTCCTGGGTTTTCCCCTTCAGCAGCAAAAGTCACTCCT CAGATTGATTCATCAAGGATAAGGAGTCACATCTGTAGCCACCCAGGATGTGGCAAGACATACTTTAAAAGTTCC CATCTGAAGGCCCACACGAGGACGCACACAGGAGAAAAGCCTTTCAGCTGTAGCTGGAAAGGTTGTGAAAGGAGG TTTGCCCGTTCTGATGAACTGTCCAGACACAGGCGAACCCACACGGGTGAGAAGAAATTTGCGTGCCCCATGTGT GACCGGCGGTTCATGAGGAGTGACCATTTGACCAAGCATGCCCGGCGCCATCTATCAGCCAAGAAGCTACCAAAC TGGCAGATGGAAGTGAGCAAGCTAAATGACATTGCTCTACCTCCAACCCCTGCTCCCACACAG**TGA**CAGACCGGA AAGTGAAGAGTCAGAACTAACTTTGGTCTCAGCGGGAGCCAGTGGTGATGTAAAAATGCTTCCACTGCAAGTCTG TGGCCCCACACGTGGGCTTAAAGCAGAAGCCCCACAGCCTGGCACGAAGGCCCCGCCTGGGTTAGGTGACTAAA AGGGCTTCGGCCACAGGCAGGTCACAGAAAGGCAGGTTTCATTTCTTATCACATAAGAGAGATGAGAAAGCTTTT ATTCCTTTGAATATTTTTTGAAGGTTTCAGATGAGGTCAACACAGGTAGCACAGATTTTGAATCTGTGTGCATAT TTGTTACTTTTGCTGTTTATACTTGAGACCAACTTTTCAATGTGATTCTTCTAAAGCACTGGTTTCAAGA ATATGGAGGCTGGAAGGAAATAAACATTACGGTACAGACATGGAGATGTAAAATGAGTTTGTATTATTACAAATA TTGTCATCTTTTTCTAGAGTTATCTTCTTTATTATTCCTAGTCTTTCCAGTCAACATCGTGGATGTAGTGATTAA ATATATCTAGAACTATCATTTTTACACTATTGTGAATATTTGGAATTGAACGACTGTATATTGCTAAGAGGGCCC GTTTAACAAATGACTGTATCTAGGCATTTCATTATGCTTTGAACTTTAGTTTGCCTGCAGTTTCTTGTGTAGATT GAAATATATTGTAAAGAAGGGATATTAAGAATCTTAGATAACTTCTTGAAAAAGATGGCTTATGTCATCAGTA AAGTACCTTTATGTTATGAGGATATAATGTGTGCTTTATTGAATTAGAAAATTAGTGACCATTATTCACAGGTGG TTCACTTTTGTTAACAGTATTTCTCTTTTATTCTGTTATATAGTGGATGATATACACAGTGGCAAAACAAAAGTA CATTGCTTAAAATATATAGTGAAAAATGTCACTATATCTTCCCATTTAACATTGTTTTTGTATATTGGGTGTAGA ACAATATTTCTTTTGTTGTACTTTATATCTTGTTTACAATAAAGAATTC

#### 559/2825 FIGURE **504**

MLNFGASLQQTAEERMEMISERPKESMYSWNKTAEKSDFEAVEALMSMSCSWKSDFKKYVENRPVTPVSDLSEEE NLLPGTPDFHTIPAFCLTPPYSPSDFEPSQVSNLMAPAPSTVHFKSLSDTAKPHIAAPFKEEEKSPVSAPKLPKA QATSVIRHTADAQLCNHQTCPMKAASILNYQNNSFRRRTHLNVEAARKNIPCAAVSPNRSKCERNTVADVDEKAS AALYDFSVPSSETVICRSQPAPVSPQQKSVLVSPPAVSAGGVPPMPVICQMVPLPANNPVVTTVVPSTPPSQPPA VCPPVVFMGTQVPKGAVMFVVPQPVVQSSKPPVVSPNGTRLSPIAPAPGFSPSAAKVTPQIDSSRIRSHICSHPG CGKTYFKSSHLKAHTRTHTGEKPFSCSWKGCERRFARSDELSRHRRTHTGEKKFACPMCDRRFMRSDHLTKHARR HLSAKKLPNWQMEVSKLNDIALPPTPAPTO

#### 560/2825 FIGURE 505

# 561/2825 FIGURE 506

 ${\tt MSAAGARGLRATYHRLLDKVELMLPEKLRPLYNHPAGPRTVFFWAPIMKWGLVCAGLADMARPAEKLSTAQSAVL}\\ {\tt MATGFIWSRYSLVIIPKNWSLFAVNFFVGAAGASQLFRIWRYNQELKAKAHK}$ 

## 562/2825 FIGURE **507**

# 563/2825 FIGURE 508

MWGDLWLLPPASANPGTGTEAEFEKAAEEVRHLKTKPSDEEMLFIYGHYKQATVGDINTERPGMLDFTGKAKWDA WNELKGTSKEDAMKAYINKVEELKKKYGI

#### 564/2825 FIGURE 509A

ACTCTGGAGTGGGAGTGGGAGCGAGCGCTTCTGCGACTCCAGTTGTGAGAGCCGCAAGGGCATGGGAATTGACGC CACTCACCGACCCCCAGTCTCAACCCTGTGAGGAAACCTCGACTTTGCCAGGTCCCCAAGGGCAGCGGG GCTCGGCGAGCGAGCGAGCGCCCAAGAGACCT CGAGTGGGAGCACCCTTCTCCGTCCCCATCCCAATCCAAGCGCTGCGGGCTTCCTGACGACGCCAAGAGACT CGAGTGGGAGTTAAAGCTTCCAGTGAGGGCAGCAGGTGTCCAGGCGGGCCTGCGGGTTCCTGTTGACGTCTTGC CCTAGGCAAAGGTCCCAGTTCCTTCTCGGAGCCGGCTGTCCCGCGCCCACTGGAAACCGCACCTCCCCGCAGCATG

CTGTCGGTGCTGGCCATGTGGGCCAGCGGCTGCTGAGGCAACGGAGGCGGCAGCTCCGGTCCGCCCC CTGGCGCGCCTACGGCGACGTTTTCCAGATCCGCCTGGGCAGCTGCCCCATAGTGGTGCTGAATGGCGAGCGC ATGCGCAACTTCTTCACGCGCCAGCCGCGCAGCCGCCAAGTCCTCGAGGGCCACGTGCTGAGCGAGGCGCGAG CTGGTGGCGCTGCTGCTGCGCGCAGCGCGGCCCTTCCTCGACCCGAGGCCGCTGACCGTCGTGGCCGTG GCCAACGTCATGAGTGCCGTGTGTTTCGGCTGCCGCTACAGCCACGACGCCCCGAGTTCCGTGAGCTGCTCAGC CACAACGAAGAGTTCGGGCGCACGGTGGGCCGGGCAGCCTGGTGGACGTGATGCCCTGGCTGCAGTACTTCCCC AACCCGGTGCGCACCGTTTTCCGCGAATTCGAGCAGCTCAACCGCAACTTCAGCAACTTCATCCTGGACAAGTTC TTGAGGCACTGCGAAAGCCTTCGGCCCGGGGCCCCCCCGCGACATGATGGACGCCTTTATCCTCTCTGCGGAA ATCTTCGGCGCCAGCCAGGACACCCTGTCCACCGCGCTGCAGTGGCTGCTCCTCTCTCACCAGGTATCCTGAT AACCTGCCCTATGTCCTGGCCTTCCTTTATGAAGCCATGCGCTTCTCCAGCTTTGTGCCTGTCACTATTCCTCAT GCCACCACTGCCAACACCTCTGTCTTGGGCTACCACATTCCCAAGGACACTGTGGTTTTTTGTCAACCAGTGGTCT GTGAATCATGACCCAGTGAAGTGGCCTAACCCGGAGAACTTTGATCCAGCTCGATTCTTGGACAAGGATGGCCTC ATCAACAAGGACCTGACCAGCAGAGTGATGATTTTTTCAGTGGGCAAAAGGCCGGTGCATTGGCGAAGAACTTTCT AAGATGCAGCTTTTTCTCTTCATCTCCATCCTGGCTCACCAGTGCGATTTCAGGGCCCAACCCAAATGAGCCTGCG AAAATGAATTTCAGTTATGGTCTAACCATTAAACCCAAGTCATTTAAAGTCAATGTCACTCTCAGAGAGTCCATG GAGCTCCTTGATAGTGCTGTCCAAAATTTACAAGCCAAGGAAACTTGCCAA**TAA**GAAGCAAGAGGCAAGCTGAAA  $\tt TTTTAGAAATATTCACATCTTCGGAGATGAGGAGTAAAATTCAGTTTTTTTCCAGTTCCTCTTTTTGTGCTGCTTC$ TCAATTAGCGTTTAAGGTGAGCATAAATCAACTGTCCATCAGGTGAGGTGTGCTCCATACCCAGCGGTTCTTCAT TACATACTGCATCTTGGTTATTTCTGAAGGTAGCATTCTTTGGAGTTAAAATGCACCATATAGACACATACACCCA AACACTTACACCAAACTACTGAATGAAGAAGTATTTTGGTAACCAGGCCATTTTTGGTGGGAATCCAAGATTGGT CTCCCATATGCAGAAATAGACAAAAGTATATTAAACAAAGTTTCAGAGTATATTGTTGAAGAGACAGAGACAAG  ${\tt TAATTTCAGTGTAAAGTGTGATTGAAGGTGATAAGGGGAAAAGATAAAGACCAGAAATTCCCTTTTCACCTTTT}$ AGGATAAATCATAAAGTCAGTTGCTCAAAAAGAAATCAATAGTTGAATTAGTGAGTATAGTGGGGTTCCATGAGT GATTTCAGTAAGTCTCATAGGTTAAAAAAAAAAGTCACCAAATAGTGTGAAATATATTACTTAACTGTCCGTAAG CAGTATATTAGTATTATCTTGTTCAGGAAAAGGTTGAATAATATATGCCTTGTGTAATATTGAAAATTGAAAAGT ACAACTAACGCAACCAAGTGTGCTAAAAATGAGCTTGATTAAATCAACCACCTATTTTTGACATGGAAATGAAGC AGGGTTTCTTTCTTCACTCAAATTTTGGCGAATCTCAAAATTAGATCCTAAGATGTGTTCTTATTTTTATAACA TCATGGCATGCTTAAATTTTAACTAAATTTTAAAGCCATTCTGATTATTGAGTTCCAGTTGAAGTTAGTGGAAAT CTGAACATTCTCCTGTGGAAGGCAGAGAAATCTAAGCTGTGTCTGCCCAATGAATAATGGAAAATGCCATGAATT  ${\tt ACCTGGATGTTCTTTTACGAGGTGACAAGAGTTGGGGACAGAACTCCCATTACAACTGACCAAGTTTCTCTTCT}$ AGATGATTTTTTGAAAGTTAACATTAATGCCTGCTTTTTGGAAAGTCAGAATCAGAAGATAGTCTTGGAAGCTGT TTGGAAAAGACAGTGGAGATGAGGTCAGTTGTGTTTTTTAAGATGGCAATTACTTTGGTAGCTGGGAAAGCATAA

### 565/2825 FIGURE **509B**

AGCTCAAATGAAATGTATGCATTCACATTTAGAAAAGTGAATTGAAGTTCAAGTTTTAAAGTTCATTGCAATTA AACTTCCAAAGAAAGTTCTACAGTGTCCTAAGTGCTAAGTGCTTATTACATTTTATTAAGCTTTTTTGGAATCTTT CAAAATTCAAAGCATGGACATTTAGAAGAAAGATGTTTGGCGTAGCAGAGTTAAATCTCAAATAGGCTATTAAAA AAGTCTACAACATAGCAGATCTGTTTTGTGGTTTGGAATATTAAAAAACTTCATGTAATTTTATTTTAAAATTTC ATAGCTGTACTTCTTGAATATAAAAAATCATGCCAGTATTTTTAAAGGCATTAGAGTCAACTACACAAAGCAGGC TTGCCCAGTACATTTAAATTTTTTGGCACTTGCCATTCCAAAATATTATGCCCCACCAAGGCTGAGACAGTGAAT TTGGGCTGCTGTAGCCTATTTTTTTAGATTGAGAAATGTGTAGCTGCAAAAATAATCATGAACCAATCTGGATGC CTCATTATGTCAACCAGGTCCAGATGTGCTATAATCTGTTTTTACGTATGTAGGCCCAGTCGTCATCAGATGCTT GCGGCAAAAGAAGCTGTGTTTATATGGAAGAAAGTAAGGTGCTTGGAGTTTACCTGGCTTATTTAATATGCTTA TAACCTAGTTAAAGAAAGGAAAAGAAAACAAAAAACGAATGAAATAACTGAATTTGGAGGCTGGAGTAATCAGA TTACTGCTTTAATCAGAAACCCTCATTGTGTTTCTACCGGAGAGAATGTATTTGCTGACAACCATTAAAGTCA GAAGTTTTACTCCAGGTTATTGCAATAAAGTATAATGTTTATTAAATGCTTCATTTGTATGTCAAAGCTTTGACT GTCCCAGAACTTAGCCTTTACCTGTGAAGTGTTACTACAGCCTTAATATTTTCCTAGTAGATCTATATTAGATCA AATAGTTGCATAGCAGTATATGTTAATTTGTGTGTTTTTTAGCTGTGACACAACTGTGTGATTAAAAGGTATACTT TAGTAGACATTTATAACTCAAGGATACCTTCTTATTTAATCTTTTCTTATTTTTGTACTTTATCATGAATGCTTT TAGTGTGTGCATAATAGCTACAGTGCATAGTTGTAGACAAAGTACATTCTGGGGAAACAACATTTATATGTAGCC TTTACTGTTTGATATACCAAATTAAAAAAAATTGTATCTCATTACTTATACTGGGACACCATTACCAAAATAAT AAAAATCACTTTCATAATCTTGAAAAAA

## 566/2825 FIGURE 510

MGTSLSPNDPWPLNPLSIQQTTLLLLLSVLATVHVGQRLLRQRRRQLRSAPPGPFAWPLIGNAAAVGQAAHLSFA RLARRYGDVFQIRLGSCPIVVLNGERAIHQALVQQGSAFADRPAFASFRVVSGGRSMAFGHYSEHWKVQRRAAHS MMRNFFTRQPRSRQVLEGHVLSEARELVALLVRGSADGAFLDPRPLTVVAVANVMSAVCFGCRYSHDDPEFRELL SHNEEFGRTVGAGSLVDVMPWLQYFPNPVRTVFREFEQLNRNFSNFILDKFLRHCESLRPGAAPRDMMDAFILSA EKKAAGDSHGGGARLDLENVPATITDIFGASQDTLSTALQWLLLLFTRYPDVQTRVQAELDQVVGRDRLPCMGDQ PNLPYVLAFLYEAMRFSSFVPVTIPHATTANTSVLGYHIPKDTVVFVNQWSVNHDPVKWPNPENFDPARFLDKDG LINKDLTSRVMIFSVGKRRCIGEELSKMQLFLFISILAHQCDFRANPNEPAKMNFSYGLTIKPKSFKVNVTLRES MELLDSAVQNLQAKETCQ

### 567/2825 FIGURE **511A**

GGCACAGGATACTGCAAATGTCCAGAAGGCTTCTTGGGGGAATATTGTCAACATCGAGACCCCTGTGAGAAGAAC  $\tt CGCTGCCAGAATGGTGGGACTTGTGTGGCCCAGGCCATGCTGGGGAAAGCCACGTGCCGATGTGCCTCAGGGTTT$ ACAGGAGAGGACTGCCAGTACTCGACATCTCATCCATGCTTTGTGTCTCGACCCTGCATGCGGCACATGC CATATGCTCAGCCGGGATACCTATGAGTGCACCTGTCAAGTCGGGTTTACAGGTAAGGAGTGCCAATGGACCGAT GCCTGCCTGTCTCATCCCTGTGCAAATGGAAGTACCTGTACCACTGTGGCCAACCAGTTCTCCTGCAAATGCCTC ACAGGCTTCACAGGGCAGAAATGTGAGACTGATGTCAATGAGTGTGACATTCCAGGACACTGCCAGCATGGTGGC ACCTGCCTCAACCTGCCTGGTTCCTACCAGTGCCAGTGCCTTCAGGGGCTTCACAGGCCAGTACTGTGACAGCCTG TATGTGCCCTGTGCACCCTCGCCTTGTGTCAATGGAGGCACCTGTCGGCAGACTGGTGACTTCACTTTTGAGTGC AACTGCCTTCCAGGTTTTGAAGGGAGCACCTGTGAGAGGAATATTGATGACTGCCCTAACCACAGGTGTCAGAAT GGAGGGGTTTGTGTGGATGGGGTCAACACTTACAACTGCCGCTGTCCCCCACAATGGACAGGACAGTTCTGCACA GAGGATGTGGATGAATGCCTGCTGCAGCCCAATGCCTGTCAAAATGGGGGCACCTGTGCCAACCGCAATGGAGGC TATGGCTGTGTATGTGTCAACGGCTGGAGTGGAGATGACTGCAGTGAGAACATTGATGATTGTGCCTTCGCCTCC TGTACTCCAGGCTCCACCTGCATCGACCGTGTGGCCTCCTTCTCTTGCATGTGCCCAGAGGGGAAGGCAGGTCTC CTGTGTCATCTGGATGATGCATCAGCAATCCTTGCCACAAGGGGGGCACTGTGTGACACCAACCCCCTAAAT GGGCAATATATTTGCACCTGCCCACAAGGCTACAAAGGGGCTGACTGCACAGAAGATGTGGATGAATGTGCCATG GCCAATAGCAATCCTTGTGAGCATGCAGGAAAATGTGTGAACACGGATGGCGCCTTCCACTGTGAGTGTCTGAAG GGTTATGCAGGACCTCGTTGTGAGATGGACATCAATGAGTGCCATTCAGACCCCTGCCAGAATGATGCTACCTGT GAATGTCAGAGCAACCCTTGTGTGAACAATGGGCAGTGTGTGGATAAAGTCAATCGTTTCCAGTGCCTGTGTCCT CCTGGTTTCACTGGGCCAGTTTGCCAGATTGATATTGATGACTGTTCCAGTACTCCGTGTCTGAATGGGGCAAAG TGTATCGATCACCCGAATGGCTATGAATGCCAGTGTGCCACAGGTTTCACTGGTGTGTTGTGTGAGGAGAACATT GACAACTGTGACCCCGATCCTTGCCACCATGGTCAGTGTCAGGATGGTATTGATTCCTACACCTGCATCTGCAAT CCCGGGTACATGGGCCCATCTGCAGTGACCAGATTGATGAATGTTACAGCAGCCCTTGCCTGAACGATGGTCGC CCAGGATTCACAGGGCAGAGATGTAACATTGACATTGATGAGTGTGCCTCCAATCCCTGTCGCAAGGGTGCAACA TGTATCAACGGTGTGAATGGTTTCCGCTGTATATGCCCCGAGGGACCCCATCACCCCAGCTGCTACTCACAGGTG AACGAATGCCTGAGCAATCCCTGCATCCATGGAAACTGTACTGGAGGTCTCAGTGGATATAAGTGTCTCTGTGAT TGTGACAATCTGGTGAATGGATACAGGTGTACTTGCAAGAAGGGCTTTAAAGGCTATAACTGCCAGGTGAATATT GATGAATGTGCCTCAAATCCATGCCTGAACCAAGGAACCTGCTTTGATGACATAAGTGGCTACACTTGCCACTGT GTGCTGCCATACACAGGCAAGAATTGTCAGACAGTATTGGCTCCCTGTTCCCCAAACCCTTGTGAGAATGCTGCT ATTGACATTGACGAGTGTATCTCCAAGCCCTGCATGAACCATGGTCTCTGCCATAACACCCAGGGCAGCTACATG TGTGAATGTCCACCAGGCTTCAGTGGTATGGACTGTGAGGAGGACATTGATGACTGCCTTGCCAATCCTTGCCAG AATGGAGGTTCCTGTATGGATGGAGTGAATACTTTCTCCTGCCTCTGCCTTCCGGGTTTCACTGGGGATAAGTGC CAGACAGACATGAATGAGTGTCTGAGTGAACCCTGTAAGAATGGAGGGACCTGCTCTGACTACGTCAACAGTTAC ACTTGCAAGTGCCAGGCAGGATTTGATGGAGTCCATTGTGAGAACATCAATGAGTGCACTGAGAGCTCCTGT TGCCTCCATGAGATCAATGAATGCAGCTCTCATCCATGCCTGAATGAGGGAACGTGTGTTGATGGCCTGGGTACC TACCGCTGCAGCTGCCCCCTGGGCTACACTGGGAAAAACTGTCAGACCCTGGTGAATCTCTGCAGTCGGTCTCCA TGTAAAAACAAAGGTACTTGCGTTCAGAAAAAAGCAGAGTCCCAGTGCCTATGTCCATCTGGATGGGCTGGTGCC TATTGTGACGTGCCCAATGTCTCTTGTGACATAGCAGCCTCCAGGAGAGGTGTGCTTGTTGAACACTTGTGCCAG CACTCAGGTGTCTGCATCAATGCTGGCAACACGCATTACTGTCAGTGCCCCCTGGGCTATACTGGGAGCTACTGT GAGGAGCAACTCGATGAGTGTGCGTCCAACCCCTGCCAGCACGGGGCCAACATGCAGTGACTTCATTGGTGGATAC AGATGCGAGTGTCCCAGGCTATCAGGGTGTCAACTGTGAGTATGAAGTGGATGAGTGCCAGAATCAGCCCTGC

### 568/2825 FIGURE **511B**

CAGAATGGAGGCACCTGTATTGACCTTGTGAACCATTTCAAGTGCTCTTGCCCACCAGGCACTCGGGGCCTACTC TGTGAAGAGAACATTGATGACTGTGCCCGGGGTCCCCATTGCCTTAATGGTGGTCAGTGCATGGATAGGATTGGA GGCTACAGTTGTCGCTGCCTTGCCTGGCTTTGCTGGGGAGCGTTGTGAGGGAGACATCAACGAGTGCCTCTCCAAC  $\tt CCCTGCAGCTCTGAGGGCAGCCTGGACTGTATACAGCTCACCAATGACTACCTGTGTGTTTTGCCGTAGTGCCTTT$ GCCAGTAACATGCCTGATGGTTTCATTTGCCGTTGTCCCCCGGGATTTTCCGGGGCAAGGTGCCAGAGCAGCTGT GGACAAGTGAAATGTAGGAAGGGGGAGCAGTGTGTGCACACCGCCTCTGGACCCCGCTGCTTCTGCCCCAGTCCC CGGGACTGCGAGTCAGGCTGTGCCAGTAGCCCCTGCCAGCACGGGGGCAGCTGCCACCCTCAGCGCCAGCCTCCT TATTACTCCTGCCAGTGTGCCCCACCATTCTCGGGTAGCCGCTGTGAACTCTACACGGCACCCCCAGCACCCCT CCTGCCACCTGTCTGAGCCAGTATTGTGCCGACAAAGCTCGGGATGGCGTCTGTGATGAGGCCTGCAACAGCCAT GCCTGCCAGTGGGATGGGGGTGACTGTTCTCTCACCATGGAGAACCCCTGGGCCAACTGCTCCTCCCCACTTCCC TGCTGGGATTATATCAACAACCAGTGTGATGAGCTGTGCAACACGGTCGAGTGCCTGTTTGACAACTTTGAATGC CAGGGGAACAGCAAGACATGCAAGTATGACAAATACTGTGCAGACCACTTCAAAGACAACCACTGTGACCAGGGG TGCAACAGTGAGGAGTGTGGGATGGGCTGGACTGTGCTGACCAACCTGAGAACCTGGCAGAAGGTACC CTGGTTATTGTGGTATTGATGCCACCTGAACAACTGCTCCAGGATGCTCGCAGCTTCTTGCGGGCACTGGGTACC CTGCTCCACACCTGCGCATTAAGCGGGACTCCCAGGGGGAACTCATGGTGTACCCCTATTATGGTGAGAAG AAAGTCTTTCTGGAAATTGACAACCGCCAGTGTGTTCAAGACTCAGACCACTGCTTCAAGAACACGGATGCAGCA ACTCCAGAACGCACTCAGCTCCTCTATCTCCTTGCTGTTGCTGTTGTCATCATTCTGTTTATTATTCTGCTGGGG AGCAATCACAAGCGTCGTGAGCCAGTGGGACAGGATGCTGTGGGGCTGAAAAATCTCTCAGTGCAAGTCTCAGAA GAAGATGAGGCCTTACTCTCAGAAGAAGATGACCCCATTGATCGACGCCCATGGACACAGCAGCACCTTGAAGCT GCAGACATCCGTAGGACACCATCGCTGGCTCTCACCCCTCCTCAGGCAGACCAGGAGGTGGATGTGTTAGATGTG AATGTCCGTGGCCCAGATGGCTGCACCCCATTGATGTTGGCTTCTCTCCGAGGAGGCAGCTCAGATTTGAGTGAT GAAGATGAAGATGCAGAGGACTCTTCTGCTAACATCATCACAGACTTGGTCTACCAGGGTGCCAGCCTCCAGGCC CAGACAGACCGGACTGGTGAGATGGCCCTGCACCTTGCAGCCCGCTACTCACGGGCTGATGCTGCCAAGCGTCTC CTGGATGCAGGTGCAGATGCCCAGGACAACATGGGCCGCTGTCCACTCCATGCTGCAGTGGCAGCTGAT CCCCTGATCCTGGCTGCCCGCCTGTGGAGGGAATGGTGGCAGAACTGATCAACTGCCAAGCGGATGTGAAT GCAGTGGATGACCATGGAAAATCTGCTCTTCACTGGGCAGCTGCTGTCAATAATGTGGAGGCAACTCTTTTGTTG TTGAAAAATGGGGCCAACCGAGACATGCAGGACAACAAGGAAGAGACACCTCTGTTTCTTGCTGCCCGGGAGGGG AGCTATGAAGCAGCCAAGATCCTGTTAGACCATTTTGCCAATCGAGACATCACAGACCATATGGATCGTCTTCCC CGGGATGTGGCTCGGGATCACATGCACCATGACATTGTGCGCCTTCTGGATGAATACAATGTGACCCCAAGCCCT ACCCCAATGGGCAAGAGTCTAGACGGCCCAGTGCCAAGAGTACCATGCCTACCTCCCTAACCTTGCCAAG ACTTTATCCCCTGTTGATTCCCTAGAATCTCCTCACACGTATGTTTCCGACACCACATCCTCTCCAATGATTACA TCCCCTGGGATCTTACAGGCCTCACCCAACCCTATGTTGGCCACTGCCGCCCCTCCTGCCCCAGTCCATGCCCAG CATGCACTATCTTTTCTAACCTTCATGAAATGCAGCCTTTGGCACATGGGGCCAGCACTGTGCTTCCCTCAGTG GTCCCAGTCCCAGCAGATTGGATGAACCGCATGGAGGTGAATGAGACCCAGTACAATGAGATGTTTGGTATGGTC CTGGCTCCAGCTGAGGGCACCCATCCTGGCATAGCTCCCCAGAGCAGGCCACCTGAAGGGAAGCACATAACCACC CCCCAGCCTCAGTCCACCTGCCCTCCAGCTGTTGCGGGCCCCCTGCCCACCATGTACCAGATTCCAGAAATGGCC CGTTTGCCCAGTGTGGCTTTCCCCACTGCCATGATGCCCCAGCAGGACGGCAGGTAGCTCAGACCATTCTCCCA GCCTATCATCCTTTCCCAGCCTCTGTGGGCAAGTACCCCACACCCCCTTCACAGCACAGTTATGCTTCCTCAAAT

### 569/2825 FIGURE 511C

GCTGCTGAGCGAACACCCAGTCACAGTGGTCACCTCCAGGGTGAGCATCCCTACCTGACACCATCCCCAGAGTCT  $\verb|CCTGACCAGTGGTCAAGTTCATCACCCCACTCTGCTTCTGACTGGTCAGATGTGACCACCAGCCCTACCCCTGGG| \\$ GGTGCTGGAGGAGGTCAGCGGGACCTGGGACACACATGTCTGAGCCACCACAACAACATGCAGGTTTATGCG TGAGAGAGTCCACCTCCAGTGTAGAGACATAACTGACTTTTGTAAATGCTGCTGAGGAACAAATGAAGGTCATCC GGGAGAGAAATGAAGAAATCTCTGGAGCCAGCTTCTAGAGGTAGGAAAGAGAAGATGTTCTTATTCAGATAATGC AAGAGAAGCAATTCGTCAGTTTCACTGGGTATCTGCAAGGCTTATTGATTATTCTAATCTAATAAGACAAGTTTG TGAAGCCCAGACATTCTTGCAGCTTGGACTGCATTTTAAGCCCTGCAGGCTTCTGCCATATCCATGAGAAGATTC TACACTAGCGTCCTGTTGGGAATTATGCCCTGGAATTCTGCCTGAATTGACCTACGCATCTCCTCCTTTGGAC ATTCTTTTGTCTTCATTTGGTGCTTTTGGTTTTGCACCTCTCCGTGATTGTAGCCCTACCAGCATGTTATAGGGC AAGACCTTTGTGCTTTTGATCATTCTGGCCCATGAAAGCAACTTTGGTCTCCTTTTCCCCTCTTGTCTTCCCGGTA TCCCTTGGAGTCTCACAAGGTTTACTTTGGTATGGTTCTCAGCACAAACCTTTCAAGTATGTTGTTTCTTTGGAA AATGGACATACTGTATTGTGTTCTCCTGCATATATCATTCCTGGAGAGAAGAAGGGGGAGAAGAATACTTTTCTTCA ACAAATTTTGGGGGCAGGAGATCCCTTCAAGAGGCTGCACCTTAATTTTTCTTGTCTGTGTGCAGGTCTTCATAT AAACTTTACCAGGAAGAAGGGTGTGAGTTTGTTGTTTTTTCTGTGTATGGGCCTGGTCAGTGTAAAGTTTTATCCT TGATAGTCTAGTTACTATGACCCTCCCCACTTTTTTAAAACCAGAAAAAGGTTTGGAATGTTGGAATGACCAAGA GACAAGTTAACTCGTGCAAGAGCCAGTTACCCACCCACAGGTCCCCCTACTTCCTGCCAAGCATTCCATTGACTG GTCTTAACTGTTGAGCCTTTCCTTTCATATCCACAGAAGACACTGTCTCAAATGTTGTACCCTTGCCATTTAGGA CTGAACTTTCCTTAGCCCAAGGGACCCAGTGACAGTTGTCTTCCGTTTGTCAGATGATCAGTCTCTACTGATTAT CTTGCTGCTTAAAGGCCTGCTCACCAATCTTTCTTTCACACCGTGTGGTCCGTGTTACTGGTATACCCAGTATGT TCTCACTGAAGACATGGACTTTATATGTTCAAGTGCAGGAATTGGAAAGTTGGACTTGTTTTCTATGATCCAAAA CAGCCCTATAAGAAGGTTGGAAAAGGAGGAACTATATAGCAGCCTTTGCTATTTTCTGCTACCATTTCTTTTCCT CTGAAGCGGCCATGACATTCCCTTTGGCAACTAACGTAGAAACTCAACAGAACATTTTCCTTTCCTAGAGTCACC TTTTAGATGATAATGGACAACTATAGACTTGCTCATTGTTCAGACTGATTGCCCCTCACCTGAATCCACTCTCTG TATTCATGCTCTTGGCAATTTCTTTGACTTTCTTTTAAGGGCAGAAGCATTTTAGTTAATTGTAGATAAAGAATA GTTTTCTTCCTCTTCTCCTTGGGCCAGTTAATAATTGGTCCATGGCTACACTGCAACTTCCGTCCAGTGCTGTGA TGCCCATGACACCTGCAAAATAAGTTCTGCCTGGGCATTTTGTAGATATTAACAGGTGAATTCCCGACTCTTTTG GTTTGAATGACAGTTCTCATTCCTTCTATGGCTGCAAGTATGCATCAGTGCTTCCCACTTACCTGATTTGTCTGT CGGTGGCCCCATATGGAAACCCTGCGTGTCTGTTGGCATAATAGTTTACAAATGGTTTTTTCAGTCCTATCCAAA TTTATTGAACCAACAAAAATAATTACTTCTGCCCTGAGATAAGCAGATTAAGTTTGTTCATTCTCTGCTTTATTC TCTCCATGTGGCAACATTCTGTCAGCCTCTTTCATAGTGTGCAAACATTTTATCATTCTAAATGGTGACTCTCTG CCCTTGGACCCATTTATTATTCACAGATGGGGAGAACCTATCTGCATGGACCTCTGTGGACCACAGCGTACCTGC CCCTTTCTGCCCTCCTGCTCCAGCCCCACTTCTGAAAGTATCAGCTACTGATCCAGCCACTGGATATTTTATATC TGGGTTTGGAGAAAGGGAATGTGAAAGCTGTCATTACAGACAACAGGTTTCAGTGATGAGGAGGACAACACTGCC TTTCAAACTTTTTACTGATCTCTTAGATTTTAAGAACTCTTGAATTGTGTGGTATCTAATAAAAGGGAAGGTAAG ATGGATAATCACTTTCTCATTTGGGTTCTGAATTGGAGACTCAGTTTTTATGAGACACATCTTTTATGCCATGTA CCTTTGGGGATAGGTTTTTTTGTTTGTTTATTTGCTTCCTCTGTTTTTGTTTAAGCATCATTTTCTTATGTGAGG TGGGGAAGGGAAAGGTATGAGGGAAAGAGAGTCTGAGAATTAAAATATTTTAGTATAAGCAATTGGCTGTGATGC TCAAATCCATTGCATCCTCTTATTGAATTTGCCAATTTGTAATTTTTGCATAATAAAGAACCAAAGGTGTAATGT TTTGTTGAGAGGTGGTTTAGGGGATTTTGGCCCTAACCAATACATTGAATGTATGATGACTATTTGGGAGGACACA TTTATGTACCCAGAGGCCCCCACTAATAAGTGGTACTATGGTTACTTCCTTGTGTACATTTCTCTTAAAAGTGAT ATTATATCTGTTTGTATGAGAAACCCAGTAACCAATAAAATGACCGCATATTCCTGACTAAACGTAGTAAGGAAA ATGCACACTTTGTTTTTACTTTTCCGTTTCATTCTAAAGGTAGTTAAGATGAAATTTATATGAAAGCATTTTTAT 

## 570/2825 FIGURE 511D

#### 571/2825 FIGURE 512

MPALRPALLWALLALWLCCATPAHALQCRDGYEPCVNEGMCVTYHNGTGYCKCPEGFLGEYCQHRDPCEKNRCQN GGTCVAQAMLGKATCRCASGFTGEDCOYSTSHPCFVSRPCLNGGTCHMLSRDTYECTCOVGFTGKECOWTDACLS HPCANGSTCTTVANQFSCKCLTGFTGQKCETDVNECDIPGHCQHGGTCLNLPGSYQCQCLQGFTGQYCDSLYVPC APSPCVNGGTCRQTGDFTFECNCLPGFEGSTCERNIDDCPNHRCQNGGVCVDGVNTYNCRCPPQWTGQFCTEDVD ECLLQPNACQNGGTCANRNGGYGCVCVNGWSGDDCSENIDDCAFASCTPGSTCIDRVASFSCMCPEGKAGLLCHL DDACISNPCHKGALCDTNPLNGQYICTCPOGYKGADCTEDVDECAMANSNPCEHAGKCVNTDGAFHCECLKGYAG  $\verb"PRCEMDINECHSDPCQNDATCLDKIGGFTCLCMPGFKGVHCELEINECQSNPCVNNGQCVDKVNRFQCLCPPGFT"$ GPVCQIDIDDCSSTPCLNGAKCIDHPNGYECQCATGFTGVLCEENIDNCDPDPCHHGQCQDGIDSYTCICNPGYM GAICSDQIDECYSSPCLNDGRCIDLVNGYQCNCQPGTSGVNCEINFDDCASNPCIHGICMDGINRYSCVCSPGFT GQRCNIDIDECASNPCRKGATCINGVNGFRCICPEGPHHPSCYSQVNECLSNPCIHGNCTGGLSGYKCLCDAGWV GINCEVDKNECLSNPCQNGGTCDNLVNGYRCTCKKGFKGYNCQVNIDECASNPCLNQGTCFDDISGYTCHCVLPY TGKNCQTVLAPCSPNPCENAAVCKESPNFESYTCLCAPGWQGQRCTIDIDECISKPCMNHGLCHNTQGSYMCECP PGFSGMDCEEDIDDCLANPCQNGGSCMDGVNTFSCLCLPGFTGDKCQTDMNECLSEPCKNGGTCSDYVNSYTCKC QAGFDGVHCENNINECTESSCFNGGTCVDGINSFSCLCPVGFTGSFCLHEINECSSHPCLNEGTCVDGLGTYRCS CPLGYTGKNCQTLVNLCSRSPCKNKGTCVQKKAESQCLCPSGWAGAYCDVPNVSCDIAASRRGVLVEHLCOHSGV CINAGNTHYCQCPLGYTGSYCEEQLDECASNPCQHGATCSDFIGGYRCECVPGYQGVNCEYEVDECQNQPCQNGG TCIDLVNHFKCSCPPGTRGLLCEENIDDCARGPHCLNGGQCMDRIGGYSCRCLPGFAGERCEGDINECLSNPCSS EGSLDCIQLTNDYLCVCRSAFTGRHCETFVDVCPOMPCLNGGTCAVASNMPDGFICRCPPGFSGARCOSSCGOVK CRKGEQCVHTASGPRCFCPSPRDCESGCASSPCQHGGSCHPQRQPPYYSCQCAPPFSGSRCELYTAPPSTPPATC LSQYCADKARDGVCDEACNSHACQWDGGDCSLTMENPWANCSSPLPCWDYINNQCDELCNTVECLFDNFECOGNS KTCKYDKYCADHFKDNHCDQGCNSEECGWDGLDCAADQPENLAEGTLVIVVLMPPEQLLQDARSFLRALGTLLHT NLRIKRDSQGELMVYPYYGEKSAAMKKQRMTRRSLPGEQEQEVAGSKVFLEIDNRQCVQDSDHCFKNTDAAAALL ASHAIQGTLSYPLVSVVSESLTPERTQLLYLLAVAVVIILFIILLGVIMAKRKRKHGSLWLPEGFTLRRDASNHK RREPVGQDAVGLKNLSVQVSEANLIGTGTSEHWVDDEGPQPKKVKAEDEALLSEEDDPIDRRPWTQQHLEAADIR RTPSLALTPPQAEQEVDVLDVNVRGPDGCTPLMLASLRGGSSDLSDEDEDAEDSSANI ITDLVYQGASLQAQTDR TGEMALHLAARYSRADAAKRLLDAGADANAQDNMGRCPLHAAVAADAQGVFQILIRNRVTDLDARMNDGTTPLIL AARLAVEGMVAELINCQADVNAVDDHGKSALHWAAAVNNVEATLLLLKNG

#### 572/2825 FIGURE 513

GAAACAAGCACTGGATTCCCATATCCCACTGCCAAAACCGCATGGTTCAGATTATCGCTATTGCAGCTTTCATCAT AATACACCTTTGCTGCCGAAACGAAGCCAGACAACAGATTTCCATCAGCAGGATCTGGGGGGCTCAAGGTTCTG CACAGGAAGCAATATAACAACAAGGTGGATGAAATCTCTCGGCGTTTAATTTGGGAAAAAAACCTGAAGTATATT AGTGAAGAGGTGGTTCAGAAGATGACTGGACTCAAAGTACCCCTGTCTCATTCCCGCAGTAATGACACCCTTTAT ATCCCAGAATGGGAAGGTAGAGCCCCAGACTCTGTCGACTATCGAAAAGGAAAGGATATGTTACTCCTGTCAAAAAT  ${\tt CAGGGTCAGTGTGGGTTCCTGTTGGGCTTTTAGCTCTGTGGGTGCCCTGGAGGGCCAACTCAAGAAGAAAACTGGC}$ AAACTCTTAAATCTGAGTCCCCAGAACCTAGTGGATTGTGTGTCTGAGAATGATGGCTGTGGAGGGGGCTACATG ACCAATGCCTTCCAATATGTGCAGAAGAACCGGGGTATTGACTCTGAAGATGCCTACCCATATGTGGGACAGGAA GAGAGTTGTATGTACAACCCAACAGGCAAGGCAGCTAAATGCAGAGGGTACAGAGAGATCCCCGAGGGGAATGAG AAAGCCCTGAAGAGGGCAGTGGCCCGAGTGGGACCTGTCTCTGTGGCCATTGATGCAAGCCTGACCTCCTTCCAG TATGGAATCCAGAAGGGAAACAAGCACTGGATAATTAAAAACAGCTGGGGAGAAAACTGGGGAAACAAAGGATAT ATCCTCATGGCTCGAAATAAGAACAACGCCTGTGGCATTGCCAACCTGGCCAGCTTCCCCAAGATG**TGA**CTCCAG  $\verb|CCAGCCAAATCCATCCTGCTCTTCCATTTCTTCCACGATGGTGCAGTGTAACGATGCACTTTGGAAGGGAGTTGG|\\$ ATCCTTCCTACTTTGCTTCTCCACCCATGACCTTTTTCACTGTGGCCATCAGGACTTTCCCTGACAGCTGTGT ACTCTTAGGCTAAGAGATGTGACTACAGCCTGCCCCTGACTGTTGTTCCCAGGGCTGATGCTGTACAGGTACAG GCTGGAGATTTTCACATAGGTTAGATTCTCATTCACGGGACTAGTTAGCTTTAAGCACCCTAGAGGACTAGGGTA ATCTGACTTCTCACTTCCTAAGTTCCCTTCTATATCCTCAAGGTAGAAATGTCTATGTTTTCTACTCCAATTCAT AAATCTATTCATAAGTCTTTGGTACAAGTTTACATGATAAAAAGAAATGTGATTTGTCTTCCCTTCTTTGCACTT TTGAAATAAAGTATTTATC

# 573/2825 FIGURE 514

MWGLKVLLLPVVSFALYPEEILDTHWELWKKTHRKQYNNKVDEISRRLIWEKNLKYISIHNLEASLGVHTYELAM NHLGDMTSEEVVQKMTGLKVPLSHSRSNDTLYIPEWEGRAPDSVDYRKKGYVTPVKNQGQCGSCWAFSSVGALEG QLKKKTGKLLNLSPQNLVDCVSENDGCGGGYMTNAFQYVQKNRGIDSEDAYPYVGQEESCMYNPTGKAAKCRGYR EIPEGNEKALKRAVARVGPVSVAIDASLTSFQFYSKGVYYDESCNSDNLNHAVLAVGYGIQKGNKHWIIKNSWGE NWGNKGYILMARNKNNACGIANLASFPKM

#### 574/2825 FIGURE 515

 $\texttt{GCGCCGCGGGGGGTCCCGGCTGCGGGGTAGACCGAGGGCC} \textbf{\underline{ATG}} \texttt{GCCCCTCTCCCGGACCCGCCGGCGTTGG}$ CGGCGCCGGAGCAGTCTACGGCTCCGGCTCTTCGGGCTTCGCCCTCGACTCGGGACTGGAGATCAAAACTCGCTC GGTGGAGCAGACGCTACTCCCGCTGGTTTCTCAGATCACCACGCTTATTAATCATAAAGATAATACCAAAAAGTC TGATAAAACTCTGCAAGCAATTCAGCGTGTAGGACAAGCTGTCAACTTGGCAGTTGGAAGATTTGTTAAAGTAGG AGAAACAATTGCAGCACTTACAGACATAACCAACTTGAACCATCTGGAATCTGATGGGCAGATCACAATTTTTAC AGTAGTCATTAAACAGATAATAACATCAAGAAATAAGGTTCTCGCAACTATGGAAAGACTAGAGAAAGTGAATAG CTTTCAAGAGTTTGTCCAAATATTCAGTCAATTTGGAAATGAAATGGTGGAGTTTGCACATCTGAGTGGAGATAG ACAAAATGATTTGAAAGATGAAAAGGAAAAAGGCAAAAATGGCAGCAGCTAGGGCAGTTCTTGAAAAGTGTACAAT GATGCTTCTCACAGCTTCAAAGACATGTCTGAGGCATCCTAACTGCGAATCAGCCCATAAAAACAAAGAAGGAGT CATTTCATCTATCAGTATTTTTACTGGAATTAAGGAATTCAAGATGAATATTGAAGCTCTTCGGGAGAATCTTTA TTTTCAGTCCAAAGAGACCTTTCTGTGACATTGGAAGTCATCTTGGAGCGTATGGAGGACTTTACTGATTCTGC CTACACCAGCCATGAGCACAGAGAACGCATCTTGGAACTGTCAACTCAGGCGAGAATGGAACTGCAGCAGTTAAT TTCTGTGTGGATTCAAGCTCAAAGCAAGAAAACAAAAAGCATCGCTGAAGAACTGGAACTCAGTATTTTGAAAAT CAGTCACAGTCTTAATGAACTTAAGAAAGAACTTCATAGTACAGCGACACAGCTGGCAGCAGATCTATTAAAATA CCATGCTGATCATGTGGTTCTAAAAGCATTAAAACTTACTGGAGTAGAAGGAAATTTAGAAGCTTTGGCTGAATA TGCCTGTAAACTCTCTGAACAGAAAGAGCAGCTTGTTGAGACCTGTCGATTGTTACGACACATATCTGGGACAGA ACCTCTGGAAATAACCTGTATACATGCAGAGGAGACATTTCAGGTGACTGGCCAACAGATAATTTCTGCTGCTGA AACATTGACATTGCATCCATCTAGTAAAATTGCTAAAGAAAACCTAGATGTATTTTGTGAAGCTTGGGAATCCCA CCTTTCACTTCCAAAGCCAATGAAGAATAATGCAAACCTGAAATCATTAAAGCCAGACAAGCCTGACTCTGAGGA GCAAGCCAAGATAGCAAAGCTTGGACTTAAGCTGGGTTTGCTCACCTCTGACGCTGACTGCGAAATTGAGAAGTG TAGAGGAGAGGGCCACTGAAAACTTCCCAGGATTTAATTCATCAACTAGAGGTTTTTGCTGCAGAGGGTTTAAA GCTTACTTCCAGTGTTCAAGCTTTTTCAAAACAGCTGAAAGACGATGACAAGCTTATGCTTCTCCTGGAAATAAA CAAGCTAATTCCTCTATGCCACCAGCTCCAGACAGTAACTAAGACTTCTTTGCAGAATAAAGTATTTCTAAAGGT TGACAAGTGTATTACGAAGACAAGATCCATGATGGCTCTCTTAGTCCAACTTCTTTCACTTTGTTATAAACTGCT GAAGAAGCTTCAGATGGAAAATAACGGATGGGTCTCAGTTACAAATAAGGACACTATGGATAGTAAAACT**TGA**GA AGCTTTTGGGGTCAGATCTCTGGAACATCATGTGATGAAGCTGACATTTTTAAAAATCAAATGATCCTTTATCTT TTCAGAAATTCATCAATTTTATAAAGAAAACAATATTGAAATTTTGCTCTATTTTCTGATCATGAAACTGATTGT AAAGCTTTTTGACAACTAATAAATGTCTTGGTAATTGCTAGATTCT

## 575/2825 FIGURE **516**

MAASPGPAGVGGAGAVYGSGSSGFALDSGLEIKTRSVEQTLLPLVSQITTLINHKDNTKKSDKTLQAIQRVGQAV
NLAVGRFVKVGEAIANENWDLKEEINIACIEAKQAGETIAALTDITNLNHLESDGQITIFTDKTGVIKAARLLLS
SVTKVLLLADRVVIKQIITSRNKVLATMERLEKVNSFQEFVQIFSQFGNEMVEFAHLSGDRQNDLKDEKKKAKMA
AARAVLEKCTMMLLTASKTCLRHPNCESAHKNKEGVFDRMKVALDKVIEIVTDCKPNGETDISSISIFTGIKEFK
MNIEALRENLYFQSKENLSVTLEVILERMEDFTDSAYTSHEHRERILELSTQARMELQQLISVWIQAQSKKTKSI
AEELELSILKISHSLNELKKELHSTATQLAADLLKYHADHVVLKALKLTGVEGNLEALAEYACKLSEQKEQLVET
CRLLRHISGTEPLEITCIHAEETFQVTGQQIISAAETLTLHPSSKIAKENLDVFCEAWESQISDMSTLLREINDV
FEGRRGEKYGYLSLPKPMKNNANLKSLKPDKPDSEEQAKIAKLGLKLGLLTSDADCEIEKWEDQENEIVQYGRNM
SSMAYSLYLFTRGEGPLKTSQDLIHQLEVFAAEGLKLTSSVQAFSKQLKDDDKLMLLLEINKLIPLCHQLQTVTK
TSLQNKVFLKVDKCITKTRSMMALLVQLLSLCYKLLKKLQMENNGWVSVTNKDTMDSKT